

cgctcgccaa ccatggaggg ggggaggaca atgccaacga ggcgagaaag ccacacgggc 120
 ccaagcatgg caaggacggc acgacccccca gaaggaataa ctacaaagcc tggcacaacg 180
 cgggcaagat accctgcaac gggcacaaag 210

<210> 29977
 <211> 426
 <212> DNA
 <213> Glycine max

<400> 29977

tgagggtggat tgcaatggag aaggtgttgt ttttgctgag gctttcatgg ccagaggctt 60
 agtgtattat tccccctctc ataatacagg tcacagatta attaattcct tcaacttttg 120
 gaatttctag atcatgaacc gaaaatccta tgaccacagc ttataaaaaa gaagagaaaa 180
 acacgtctaa ttataattat tatattcaat tataagttta tttatcgatt aagtatgata 240
 atcattatga agatattcaa atgatgacct actgggtttc tgttaataata acagaggacg 300
 catcaagtat ctaatacaag tgttatgggc tgggtgtgtca acgcacaaag cttgtgtata 360
 ggaagcttat ggtggtataa tccccctgcag aaatacaaat attaattatt ttaacagatt 420
 tgtctc 426

<210> 29978
 <211> 142
 <212> DNA
 <213> Glycine max

<400> 29978

tatccgaacc tacctactca tactttatgc ctagactgat cggctactct gcccttaat 60
 ctttctatgc atagagcata ctgtcaatga gacagccaag taccaactaa tctcaagaga 120
 aacatgtcat caagcttcat at 142

<210> 29979
 <211> 355
 <212> DNA
 <213> Glycine max

<400> 29979

ttacaatcgt tgaccatgat ttgagaaacc gtgagggtta atatcactgt tgctatgaca 60

atcttttgtt atagtcagtc cacatggtag ttatgtcaca gggaatgttg actataaata 120
tcacgtctaa gctatcggaa ctgactccga acttggatac tgactatagt gattcaacag 180
actttttgag actacgggat ctatgagaga ctacagcatgt attctataat gactatacaa 240
agtattttcg cactataacg actattaata tcatactcct gagactatgt atagagtaga 300
tgagttatat taagctatgg atcggacatc atcaagtata ctgcttgatt atata 355

<210> 29980
<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29980

agcttgatnc tcgaatngat tntagcctta gtttcacttt gggtattagt caattcgggt 60
aagaaagaaa aatcccanag aanaacgtcc gatttgattt ttgattatt ttattaaaat 120
atatattttt tattattata ttactatttt gccttttttt tgttttaaata gtgggtacgg 180
catgacagaa cggtcgaatt tcattttaac agaaattaaa agatgttaca attcaaatga 240
tcgggtgaaa tttattttat ttttgattag gcgaggaaat gacttanata aatgactaaa 300
gcacgtcaaa agaggggatg gaaagtaaata gaaataaaaa taaaagcacg cgaaacaaat 360
ggggaccact aagggtacat agaatg 386

<210> 29981
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29981

nttaactgaa tttgcaacgt tctaattgtt ttttaaatgg tgtaatcgat tacaatatat 60
tggtaatcga ttaccagtgt atctgaacgt tgaaattcaa attcaattgt gaagagtcac 120
atcttttcat aaaattattt gtgtaattga ttacatgggt ttggtaatcg attaccagtg 180
acaagttttg aataaaaaatc aagagatgta actcttccaa tggttttcag gttttttctca 240
aggttataac tcttccaatg gtttttcttg accagacatg aggagtctat aaaagcaaga 300
ccttgacttg aatttcaata actntatata tatactttta catcctttga atctctttga 360

acatcttttt gaactttctt ttctttctct tcctttgcca aaagctttct gagttttctg 420
gtttccaaac cttgttcttt 440

<210> 29982
<211> 371
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29982

agcttgacag gtccaggtgc ggttgctgct actggtggag ggacttcaat ttgcttgcca 60
gacctcaagg tgatggcact cacattcttc ggattntgca ccatttgtga aggcaatttg 120
tcagaatttt gggactgagc ttggttcaac tgagtagcca tctgccccat ctgatttctc 180
agactctaaa tagaggctct tgtctctntc tgaaattgca tattctggat agtcatttgc 240
ctcactaact cctctaagga aggttgagaa aagggcctca gttgcttggt gtctttgttg 300
gtggtgctgc attggaggag gaacatatgg cctgcttgga ccaacaacat tctggaaggg 360
agggacaggc t 371

<210> 29983
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29983

tcctttctt tggccaatgc tggactcggt tggcagtgat ttccttgcca atctgatgct 60
cagaaacatc aatatccacc actccttcag taggtctgcc caggtatttg ttgatcaccg 120
caggggagaa tctaacacac tttcctctga caaacactct ttgataatca tcactttttc 180
tgtttggtat gtcagagggg atgttgacaa taaattccct gactaggctt tcataacagt 240
ctcccaactt ggtgactggt ttcagtagtc cagcagcctt gatgagttcc atggtctcct 300
tgcaatccaa ggcatttctt ccagttctc tttccaaggc aagtctgcgt tgatacacgt 360
atttccacct ttcagcattg ccaatggagt ggaatgagat gttgtccaat ggtgcatcan 420
ggacattntc aggcaccttt ttcccagatt tcttggt 457

<210> 29984

<211> 462
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29984

ctaagctgag tcacattggg ggttccagtt aaacctgcgc tgtgtatata cctcattatc 60
 aattggccta gtatcaacaa cgttatcaac tgtttgaatg ccctccaaaa gagttgggtg 120
 anaactatca ttacttgaag gcagatattt atacacgact aataatccaa gagtgaatat 180
 agcttgcagc ttcgtagagt ccccttgaac tgagaagaat atgagcatgg cacgtatgaa 240
 cagatccaca cagagacgca tgatgcatat tacagtgtct attaacctcc catcgttgcc 300
 tgcggggaag ccacgaactc gatacacaaa cagagtattg tacttgtcaa ccacatatct 360
 atacccaaaa taaatggcac caacaggaac cacaatggga ttaaatgaac agtatactag 420
 agtcagggct aatattgtca aattaaaggc gtaatactgt gc 462

<210> 29985
 <211> 411
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29985

agctntggcg gtcnttggga ttttctcaag cccaaccca tcaatgccta agacaaatcc 60
 ttttgggggt tcatttaata atgtagcaaa caaagttacc aagctctaca gttacattct 120
 ctgttcctag tggagagttg tttaaaacca gcattcttta atttctcaag ccncagtc 180
 tcttcaccaa ccagacaac aatttcaggt gcattctctg gtggcttcaa tgctgtggca 240
 gctgttcccg cacaagctcc aggtgggttt gggcagccag cacanattgg atcagggcaa 300
 caggttcttg ggtcagttct cggtggttnt ggacaatcaa gacagcttgg tagtgggttt 360
 gctgctccta gtggtttcgg tgggtgattt gctggcggtg gttctccag t 411

<210> 29986
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29986

tgtcactttt gggaggattg taccaaacca tcacccccag atcaatcttg actgatctaa 60
 aaccccatth atgctagtgt actttttattc agtcactttg gaaataattt atagaaatac 120
 tgtgccatat atttttgaaa caaacattat aataaactac ccacatactg actntaaaca 180
 agtacttatac atttagttaa ttcatacctaa atacatggat atgtccaaaa tcctacttga 240
 agcaccaatc tgacattata aacaatggcc gccatctaac actagcattc acttgtgact 300
 ctagatattt ccagcaagtt cattgttcaa aatcttaatt aattgcgaga ttctgcccta 360
 ttttgcatag ataaaaagta atataatata gattattntt tacttanaca agtagacatc 420
 taggatttgc aatanagata gagcaactaa c 451

<210> 29987
 <211> 427
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 29987

agctttcagg tttanactna cctaagcaca gtcttagaaa cttacatag gtcctgtcct 60
 tgaaagcaag ctttctaaga catttctaag acagtgcata cgtaagcact gtccttgaaa 120
 gcaagctttc taagacgatg cttacgtagg cacagtcttg gaaagcaaac attctaagac 180
 ggtgggttacg taagcatgtc ttagaaagct tactttctga gacgggtacct acaaattacc 240
 gacttcgaaa gttggctatt ttccaagacg atgtgttctt actcgtcgtt gaaaggtaac 300
 actttcaatg gtgttagctt ctacgacggt cgacaatcgt ctttgtatat taatttggac 360
 cgtcgtagaa aaaccatttt tcagtagtgg ttttaaggaag ggtgaaactt aattccaact 420
 cattcac 427

<210> 29988
 <211> 459
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 29988

tctactgagc acaaataaaa acaaagcaaa atttattttc aatcctacaa aaagaacat 60
 aaattgggga aaatatatac attttgtaaa agttttctat acaaaaagta gtcgtataag 120

acgactaaca acatccatca aatgaaaatt caaaaaccaa aaggaataaa caaatcaagc 180
atgtcagagc aaatttatta ttatattcaa caagatcgta aacattggag agtaccgtta 240
gtggtggtag tccaccggag tgggcagcag ggaataaaag tgccttcgc ttgtcatcat 300
gcgttaggtc tcatcttcgt gcaatcaatt ntgatttcaa cttgaatgtg aacaagcgcg 360
tttatgtgtg tagtggtgag tgagagggga agtggaacat gtgagtttct ttntatttaa 420
gtggaaaata ttctaagacg gttatatggg aaccatctc 459

<210> 29989
<211> 193
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29989

tccttacgca tctgtgcggt atttcacacc gcataatggtg cactctcagt acaatctgct 60
ctgatgccgc atagttaagc cagccccgac acccgccaac acccgctgac gcgaaccctt 120
tgccggcgat gagaatatga ccantggtgt tgatgcacta ttacatgcc ctttgactta 180
tgacttgatc gcg 193

<210> 29990
<211> 420
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29990

cgtaatccta tgaagctgtg cacaatcnga cgacntcagc tcggaccggt gatcctcaga 60
gtcacctgcg gcatgttgct ncatatttcg catataanga acataaagct tgggatcgat 120
cgggcccccg gtaaaggagg ccatggaatg gctcaatttg gtatgggacc aatgaaactt 180
ttttgattta agtcataaaa tgatgccaaag gtctgttgat ccgctgaaac ttatgacatg 240
gctgatcaca agaattataa atgaagctac cctggattca aaagaatcca agagttgcat 300
gaagacacat caagtccctc tatacaatgc ttggcattt cagcactctg tcaatagttt 360
ctttgaatgg caggtaacct gnggttattc tatttattaa ataaatataa ttaattaaat 420

<210> 29991
 <211> 445
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29991

tggtttatga attgattnta gccttagttt cactttgggtt attattcaat ntgattaaga 60
 aagaanatcc caaagaaaaa tgtccgattg gattttttttt attattttat ttaaagatat 120
 ttttttgatt attatattat tattttgcct ctttttgggtt ttaaactatg ttacagcatg 180
 aaagatcggc cagattntat tctaacagaa attaaaagac gttaaaactc aagtgatcag 240
 tggaaattta ttttattttt tgattaggcg agaaagtgc ttaaataat gactaaagca 300
 caccanaagg tggtagagaa agcaaatgat atanaataa aagcacgcga aacaagtggg 360
 gaccactaag ggtacataga atgaattgaa tggttcgatt tcggaaactt accggttgaa 420
 gaccaaacia cgacgaagaa cgatg 445

<210> 29992
 <211> 409
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29992

tcgcttacca attgtggaag tgaagctctt ccatgccttt ctagcaggtg gaaggcttag 60
 ccttagcctt aggcttatgc aggatttcat tgataggagg agagagcaat gcttcaaata 120
 agaagaggag gatttaatat gtgggttaaa gttaaaactca agttcaaata gaaacttggg 180
 tccttaattt ataacatggt agcatgggtt agattccctt ttttgcagtg tcatgttgcg 240
 gaatagtata ttactgtatt catatatggt gggtttgctg tgaggggttg cttagttatt 300
 acgcgctcaa ttttttttga caagacgagc atanagctgc agtaggtaaa atctgccaaa 360
 ttctccagca ccaacgacct agcttgctta attattaagc tgacattaa 409

<210> 29993
 <211> 450
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 29993

caattatcta cattatccac attcctagct ttctttatcc acatctacaa aatagtcctt 60
agatttcctt ctctctttct tttgtataac tgtgtcatat cttagatctc ctttcacttc 120
cttcaaacia caaaatcatg ggtctgagag acattgggtgc ttcactgcct cctgtgtttc 180
ggttttatcc gagtcatgag gaattgggtc gccattacct ctacaaaaag atcgcaaagt 240
aggaagttct gaagggtacc ttggctgata ttgacctcca catatgcgaa ccttggcaac 300
ttccaggtaa atatatattc attctaaatt atatatatat atatatatat atatatatat 360
atatatatat atatatatat atatatatat atatatattc ataagctatt ctgaattata 420
caccttecta tagcttgatc tctgtggttn 450

<210> 29994

<211> 450

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 29994

tgtaggatta tgggggtacc atcacatgtg gtactaggtg gcggtcgggc gatggtgcac 60
aacaagcttt ccacatccac aatgcgcgca taaaccaccc atccctgtt gccacacctc 120
aacggagctc acgtactccc acgtagccca tatcctcggt tctctcaaca ccgggtcccc 180
atcaatcctc tcaagcttcc acaacatcca agcaaaaacia cattcaaaca gcacaagcta 240
tcacagccaa gcaaaacaga gcaaaggcag aaaactctgc tcaacacatc aaccaaaatc 300
acagcttttc tcatgtaaag accacagtaa caattccttc gatccaattc gttaaccggt 360
ggatcgactc caaaatttta ctggaagtct atagtgtata agcctacatt ntgaccggtg 420
ggatatacta gcanacatcc agaacgcatt 450

<210> 29995

<211> 304

<212> DNA

<213> Glycine max

<400> 29995

tcttgaacga accccttaaa tgcaaagcca accatctctg tccccatcca ttcatttctg 60
ccgccgaaca ggtgatctgg atcttgacct ccttcccttg cataaaaagg ttacagtaaa 120

taaacaacgc accattgtaa cataaatatg aagcctgaca aattcacctg tgcagcttaa 180
tcttatatga tgatggaagt ttgtctgaaa aacataaatc ttaatcttga aaaactctgg 240
caacaaacct ggaagaatca agagaagtat atggagaatg cctgcgctga gaatgcttgc 300
tatg 304

<210> 29996
<211> 413
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29996

taggattaaa gttcagaaac ttaagagccc aacaagcttt atgctctagc tctatgggta 60
ggtgacaatt ttatcaaaca caagtctata aggagacata ccaatgagag ttttaaagt 120
tgtcctataa gcccaaagtg catcatctag tttaattgcc caatctttcc tagatgcact 180
aactgttttt tcaagaatta tttttaactc gctattggac aattctattt ggccactagt 240
ttgtgggtga tatgggggtg caagcttttg agtcacccat atttagccaa gaggccatca 300
tacaacttat tacagaagtc agtgcctttg tcaactaatga tngctcgagg tgtgctaaat 360
ctgggtgaann atattttctt tgaaactttt ttaccaccag agaatcatta atg 413

<210> 29997
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29997

tttgcattgca gcttatctat tctctttnat aaaaagaggg gaaataagaa aagtgacatg 60
ggatgaaatt cttttatatt tccatgtttg taatacttcc tttccaatgg aaacgtcgga 120
aatccaactt atgaagagaa aggggtggtc gtagaactgt cattatgcat atcatggtag 180
ttacttatga taatttgagt gatgcaagtc gtcttttatg tatattgata gaatacaaac 240
tctattgatg agaatggaag agtatagcag tcaggacaag acatcctcga acagtagata 300
gaagaacaaa ctatagaata caatagaaga tataccagag attgacatat gtacattcaa 360
accatctaga agtaatgctc tttgaagagg acaga 395

<210> 29998
 <211> 440
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 29998

tattgtcatt tttcttttat tntcttctct cataagtatt tatggagaga tttatccaaa 60
 caaaaatacc atgaagaaca taaaattagt cagcttaacc tcaaaagtac aaagaagggt 120
 aagtcaattc ataaactatc attatgttgt aacacttaca atttggaatt catctaattt 180
 gctcctgaa caagattgga aatgaaagt agcacaaca gtaacgaata ggtcttgcta 240
 ctatagtgga gagaggaaat tctgatggtc ccacaaatat aacaagatat aaatattaga 300
 taaattactt gttccttaat ttatttactg tgataacaat tgaggttggg tgcattgagca 360
 atggaagaga aggaactaat aatttcttaa ttatatgttg agcaaaatct attaaataaa 420
 aatggacatc aaaagtaaat 440

<210> 29999
 <211> 399
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 29999

agcttagtcc tcgtccacca gngacctgaa ttgaatataa gaacatcaga atcaatccac 60
 tcgtggctaa tatcatcaat cttgtccaat ctcaagtgcag tcttcaccct ttggggagca 120
 tgtctaggca cagaaccagg cctcaccagg aacactgac gataaaaatc gattctaaca 180
 tcaaaagtac taaacctcac acctanaac cttatctgtt ttgtgatctt gttgcctttg 240
 atttcataga cactcttctt atcctcaact cctgtcatta gcaaacatat catggactcc 300
 cattgcgtcc tactcaaaga atcgccgacg aanaccaccc ttttccacg aagtcgttcg 360
 agtattccac ggacatcaaa ccttggaatc tcacagttc 399

<210> 30000
 <211> 442
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30000

tcttccataa accataattc cctcctctgc atcctgcaca cattcttgcc gacttttttt 60
 caaactcata tatatgttgg gaaatcacac aattgcccc gtccctcagc tagctagctg 120
 gatttgtaa ttagtgcta gctgaattta accaagattg tgaattagca ataataagtt 180
 aataactcat caaatctcta tcaacttacc ctatatacat aaacgtctct gttttttaaa 240
 cttttaaaat aactccaagt caattcatat tcttttagact ttagtgggtca cactcacaca 300
 ctaccaatat aaataatttt acaatctttt gaaataaata atataaaaat catataggta 360
 gtatatgatc tattcattta ttntttcttt atcatactta taaagaagcc atgcatcaca 420
 nagtaacccc atggaagtgg cg 442

<210> 30001
 <211> 101
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30001

agcttggcca gtagaangaa gactcgacac tatgtagtgt gggtttgaac cggaatggc 60
 tgcccattgg agaagaatga aatcaatgag gagttatgat g 101

<210> 30002
 <211> 448
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30002

acgggtcatt ntgaccagag ctccaaagat ccataatcag cagaaaattc tgaaaggaga 60
 aattacaaaa attgaagata aaagctggga aatcaagag taagatattt ttaaaggata 120
 aatcagctca agaagaacat aattacttga attaattaga aatgctattt atttttaatt 180
 tcttgtagct atctccttaa ttaaactctat ctacaattct aaaaataggg ggtagacat 240
 tcgttgtaac ggtgtaaaag tccctatcaa ttcttagaat tatatttttag aattgcacct 300
 actacatgtc tccattattc cattagacac tctagatttc attgtattat ttttataagt 360

gcaattcctt ccacctagga ggagaaatga tgaacctttt ttcgctgtaa ttctatcaat 420
tcaatacttc atcttaagtt ctttattt 448

<210> 30003
<211> 411
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30003

agctngnatc gttganaang cctcttggca attctcagca gagaaatgga ttttgtcaga 60
ttgagacgaa acagttatgg ggaaatcagg tagctgcacc ttgaagtgtt ctccctcgat 120
ccttgacaca aaggacatta ctgacccaac ccaggaaatg gggaatactc atccctgtgt 180
ctggacttag tcgtagacta ctaataaata gctttcagcg gaaagtttct tttgatgaat 240
aattggaacg ttataattct ctgaaatctg ttatattgga ccagagtgga caaaatggca 300
ttgtatgaca gtcacttcct gagacaacaa cctncanaga gaaagttggt tctggatata 360
ctataaagtc acttcctcct tcacctacac ttgaacatat gaaaatatcc t 411

<210> 30004
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30004

tcaccttgaa gctaattgaa ataagctaan aaaagtcata actttaaaat aagctcttac 60
aaacacctca tatattcaaa acgagaagaa ctgtggaatt aaacatcatc cactctatgt 120
cagactggta attccgtaat tgctgcaaact actaagttat tcctctagca tccattgtcg 180
atttcaagat tttcctatca tagtgatcac attatctgat tgtagttatc attggtacac 240
tntaacctca taacaaaagtc aatcaataag agtcgatttc aatttgcaga tcagaaaagc 300
taccctcaa attaggttat tgacttaagc agctgaatgc ataatacaaaa ggaatttgta 360
gaaactaaag caaaatatat agatggataa tagaaaaagg aaaaggctgc acatctcaat 420
tccttaccga ttcaccactg agattcatgt aag 453

<210> 30005

<211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30005

gagcttggct ggagntgagg tacgtgtctc ttgaaattg aagcgtctgc aattgccata 60
 tctgcatga cataatggcc atccaaactg cgcaaaact atatatcaga tgttgaatca 120
 tgcataagat atttcagcat gtctactata tatactatta tgagcattac aagagcaact 180
 gaatgcatca cgcgtgcata tatacgtgtt caataacgtg tacgcctaac tggcgaagat 240
 acatataggg cacttggcaa ctgaatcatg atgatacgag tacctcagac aaaacactcc 300
 tcaacaggaa tatgaattca cagtgtcaat aatgctgttg gtcacgaga taactcctct 360
 tccctctccg ctctttcgtc cacgatatat attatcaaaa ggctaataatc attgat 416

<210> 30006
 <211> 377
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30006

agcttctcag tgttgtntct ctttttgcta cctangctat aaatagaagt atgtgtaaca 60
 cttgtggtaa ctttgatgga tgagagtctt gtgagacaaa cttcaaagtt caacttctct 120
 ccctcttttc ttccttcaat gtcgtgtctt gccgtctctc tttcttttct tccattgaag 180
 caccttctcc aagcttctta tccaaggcat tctcttggtg gcgaagctcc ttcttccatg 240
 gcttattccc tagtggatgg cgctctgtct cacctcttct cctttatctt ccgctgcac 300
 tccatggtag aaaataacca ttgaaggacc tcaatgaagc tcaaagatcc agcctccata 360
 gaagctctac aagtaag 377

<210> 30007
 <211> 454
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30007

ctcaagcttg tagcatcgg actaaggttt gagagctttg ggcacccggt cctatgggtg 60

ctgacggagg agctggagtg gatgatgaag ggatgtcttt tgctctagcc ctttttcttg 120
atgacatctg taactaaaaa gaactcaaaa ttccttagac caaattaacg atgggcgctt 180
agcgggatac aactcgtca gtgcgcctc agaaatataa catatcggct tagcgaaaca 240
gcatgtgctt tagcctaatac aacgctgcaa cagatatgcg ctaagctcag cagggttgcg 300
cttagcggca gcatgaaatt cagaaaattc actaagtatg ggggcttagc gagcaaggct 360
cgtttagccc aatggctgcc acaatgaaat gagcttagcc cagataggct cggcttagcg 420
catagctntc aacaaaaaat tggactaagt tacc 454

<210> 30008
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30008

ttgatctcct tcttcagcct ctggatctcg tccttcttta gccttgcaac gctcatctcc 60
atcgaagctt gatgacgaga ggccagagag tgtttcttct ctctgccccg ctctctatga 120
tggcaacgaa gagtcacgat acaagacaaa gtaagataga gggttgtcaa agcttatcca 180
gttataacgt gccacgtag catacaaatt ggaaaacaca ggttttcaaa gacggttttt 240
taaaacgct ctctgaagc atattttaag ccgggtgtaa agtaccgctc ttacaaaagc 300
tataattatg cacaaaaatg tcaccgctnt atatactaca tcggttgctg tataaccgac 360
gtataaacag tgacgtagaa aatctctttt ctagta 396

<210> 30009
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30009

tctctaagct tctcatccga gacactctct tgggtggtgaa acttcttctt caatggctta 60
ttccctaata gatggtgtct catctcacct tttctccttt atcttacgtt acaacttcat 120
ggctgaaaat caccattgaa ggacctcact gaagctcaaa gattcagcct ccatagaatc 180
ttctcaagca agcttccatc aaaaagtact gaacacaact tgctatgttc aaaacagaat 240

tggtccaagg aggtgcatat ttgtacacct

450

<210> 30012
<211> 482
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30012

ncgtgaatcc tgctgatgct actgcaactt cgggcgaact gagctcgta cccggatcct 60
ctgagtcgaa cccgaggcat gcttggcaac gaggtcaatg tcttatccgn ggangccatt 120
cttagaagct attaaatggt ttcaggcttg agaattaact ttggccagaa ccactttggg 180
gccattggcc catctgaaaa atgggtggtgg gccgctgctg aatatcttaa attggccatg 240
cctcaattcc ccttttggtg cctaggggtg cctataagca ttaatctgag aaaaatatgg 300
tgtgggagcc tatcattaaa acggtcgagg ctaagttgaa caagtggaat caaggaacat 360
ctctatggct ggaagaatca cccttatcaa tgctgtttaa cagcacttcc cttgtttact 420
tgtctnttac aggtccctc agcagcatta atagataaat gctattcgag acacttttgt 480
gg 482

<210> 30013
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30013

tagtgaaaca ctaaataac acctctcttc taatggaaat tcaaacaatca atttgccaca 60
taatgtaata ggggtgtaaa acaggttgaa ttcacaaac caaccacat aatcctcaa 120
aaaatggcag agtatattag gtgattaaat tcatcatccc accttgata agttgtcaac 180
ttaaggagtt ggaggctaaa gaactaactt gctcagtggt atcaagacta taccgatcat 240
gattcggcca ggggaactagg ttgatgggtc agtggttgaa ctgaacgtgg gtcgctagcc 300
gaactggtat atattaaata tttaaattct atagtaaata tatcatatat aaattacttt 360
ntttgtatct atatcataaa attntgtttg tatcaagagc gggtggcaca attggtagcg 420
gcttaagtcc ctaaaccaag tggtccaag 449

<210> 30014
 <211> 367
 <212> DNA
 <213> Glycine max

<400> 30014

catcatacca cctccaggtg ctggaactac ttcacatggt tcttgatggg gcctatgcca 60
 attaagagcc ttggatgaaa gaggtatgcc tatgtcttcc acaacagagt cacacttaga 120
 agacggactc ctacctctcc gtattaaatc atgaaagga agaagccaac tgtcaagctc 180
 tttcacatct ttgagaagtc cctgttacac tttggcgaat caagatctaa tgaaaaagat 240
 ggatcccaac agtgaagctg gactattcct gggatactct acctacagca gagcatatag 300
 agtatacaat tccataacca tagcagcgat ggaatccatc aatgtggttg ctgatgatct 360
 gtctcca 367

<210> 30015
 <211> 291
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30015

tagcnnagnn gttcctgcct catttccagn gagcatcgcg atcaaacaaa gagcgcggtat 60
 atgacaatca ccgggtcaat aaacgaagag gacaacactg tttaccttaa agaccggata 120
 tctgacataa tgggtatgca caatttcac cccacattttg caagaatctt acaataaatt 180
 atttcatgga tcaaaccacg tagttatgtg ccgggtcaatc tccacatgat gcataattaa 240
 gctattttaga ttacggagaa gagctgataa aaattgatta ctgatcacct c 291

<210> 30016
 <211> 445
 <212> DNA
 <213> Glycine max

<400> 30016

taatggcctc attagtctca gtcttaggtc caacagcttg aggtggaacc ttccaactga 60
 cctactttcc cttccttccc tgcgctttgt atatctccaa cacaacaact tctcaggtgt 120
 tattcctgat tctctgccac cagggcttat ctttcttgat ttgtcccata actcttttac 180

aggacaaatt ccagcctcaa tccaaaactt gacacatctg ataggattta acctccaaaa 240
 caactctctc acaggaccta ttcctgatgt taaccttcct agccttaagg atttggattt 300
 gagcttcaac tacttgaatg gatctattcc ttcaggtctc cataagtttc ctgcctcctc 360
 atttagaggg aatttgatgt tatgtggagc accttgaaa caatgttctt cagtttcccc 420
 taataccaca ttgtctccac caaca 445

<210> 30017
 <211> 310
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30017

agcttcntag ttaaattggac ggaccttann agaatagctt taatagccct tgtgagcctt 60
 gtttccctta tcttgttttg aagctcacta caagccttaa gtgagaaacc atgatatcac 120
 catatcctta aggaatattg gagctgtgga attgttatgg gaataagagt ggagggtttt 180
 tgtttcattg gacaacttgt tatgatggct atgctacatg atgtattttg tgccatactt 240
 gatgtacatt gtatatngga taaatgttgg acatgctgaa tgaaatgttg tgtctcaaag 300
 gctatagagt 310

<210> 30018
 <211> 430
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30018

tgtagaatgg ctagacatga tacatgtcan ggtttggttt ggttcaagga taaaagggat 60
 gccccacatt atttccatga cacaaatgca aaaatgatga tttggaaatt ttatgcaaaa 120
 ctggtcatgc atgcacctat gtggacgctc aagtgtcaaa cttttatggt catgtgatgc 180
 tagggctcaa gattcatttc ctctatttta aatcaacca atgtttccaa aatatgttct 240
 tttatcaatt tgtgcattca tccgagtcca tttcgggcgt ctgggaaaat ttcacagcgt 300
 tcacccttcg ggtgtacaca cacatttttt tcaaaactag ctatgatcag cgaatttttc 360
 ttcaaagaaa agatggaagt catctctttt caaaagcatg ttggcttgtc agctatacta 420

cttattattt

430

<210> 30019
<211> 424
<212> DNA
<213> Glycine max

<400> 30019

taagctgccg cctaagattg ttatatattgc gtggaggcta acttttagatc gactaccaac 60
tagagcaaac ttgcggtctc gacagatcga agttgaagat gcaacatgcc cattctgcag 120
agaggtggat gaaagcgcg gtcacatctatt ctttcattgt cacaagataa ctccagtctg 180
gtgggaatcc ttgtactgtg tagatctttc cggtgccttg ccaaatacacc caaggcatca 240
ctttcttcaa tacatacaca gagtaacaga ggaaatgacg tctaccacat ggaaatgggtg 300
gtgggttgga ctgacatgga ccatttgga tcaaacatat aacattatct tctccaatgg 360
tacattcaat gccatcgaga tactagatga tgcagctttc ttactatgga tgtggctaac 420
taac 424

<210> 30020
<211> 390
<212> DNA
<213> Glycine max

<400> 30020

atcttgaagc tatatagcgt ggaagagtca atcttcttac ttttatttgt tgaccacaaa 60
gtggtaccta gagatatgtc gcgggggtca ggagaccttg gggacgtcag gtggtgtgct 120
attgccc aaa accaagcttg atcaatctg acccaactcg ggcatagtca gtcagtgaga 180
acctgtgacg tacctaaaca ggcaagctcc tgacagtcaa ccaataaaaag aacaaagacc 240
acaaagcaag gaggcttgtg tgggtggctgg ccagctatgg atcttgagtg gtatctggaa 300
tttggcctct ggtaatcgat taccaagggt gtgtaatcga ttacaaggct taaaaatgaa 360
gacaggaagt taagatggcc tctggtaatc 390

<210> 30021
<211> 439
<212> DNA
<213> Glycine max

<400> 30021

tgtgggttttc tcacagatag gacatgcatg atgccttttc aactgtatc cacttaaatt 60

tccatgatgt ggaaaatcgt taatagtaca aaacaccatg gcgcgtaacc tgaacatctg 120

ttgcacattt gcatcccacg catctaccct ttcttccac aattttttca aatcttcaat 180

taacggacta agatacacat caatatcatt ctcggttgct cttggaccgc cgatcatcat 240

acacaggata atgtattttt gcaaaatata caaccagggg ggaggttgta aatcatcagt 300

aaaacaggcc acaaactgtg gttgttgctt aagctgccat aaggattcat tccatcagaa 360

gcaagagcaa gccttaagtt ccttggtctg tccccaaact ctggatacaa atgatcaatt 420

gtcttccact gtggagaat 439

<210> 30022

<211> 402

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30022

agcttcttat ttcttgctca tcttggttgt gatactntnt cttccatggc ttattcctta 60

atggatggtg cctcctctca cctcttttcc tttgtcttcc gctgcatctc catggtggaa 120

aatcaccatt aaaggacccc attgaagctc anagatccag cctccataga atccctcttt 180

gtaaacaacc aaaatttctc aattgattat ttttccttgt ttggtgattg ttgcaattct 240

cttagtgtag tactagtga atgaaatagt gtgttaatct ctctctcca tttctctagt 300

ttttattttc gacttgaatc ctttacgaac cctattctac aagttgttga actatattcc 360

aaatttctac cttttgcaac tatggaacan taaaatatta aa 402

<210> 30023

<211> 459

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30023

tccttgagaa gatntataga gaagttagag cttaactaaa cacaccctc taatagttaa 60

gtcacctcc ttgagatgag aagctagagc ttaactacac acaactccta taatagctaa 120

attcatccca tgccaaaata catgaaaata caaaaaaatt tctagtacaa ggactactca 180
 aaatgtcctg aaatacaagg ctaaaatcct atactattag aatgaccaa atacaaggct 240
 caaaagaaga aaaaatctat tctaattttt acaaagaaga gtggacccaa cattgacca 300
 tgagctcaaa aatctatcct gaggttcctg agaaccccag agccttcttt agcagctcta 360
 acccaatcat cttggagtct tctgtccaat aaccttgga gaaaggattg catcaacttc 420
 tccctcgagc gttnttgat ctaattatgg tgtaagagt 459

<210> 30024
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30024

agctntaagt ctttcaacan attatttatt gttcatagtt ttattaataa aaaaaataaa 60
 tgccttatat tctggcaaaa aaggtacata tgttgacaaa acatatgtan gaatacaagt 120
 gggaagggaa gtctcttttg tgtaagaatg aaaaagtgag caccatatga gtgaggataa 180
 aaatcataaa ctngagtttt aaagggttaa gttaaagtgt gaccgtcaat tttcttatgt 240
 ggntgttcat agcttatagg taaaatctcc cctgtgattt atccccctca tngcataata 300
 attagtataa gagtagatga tntaacttgg tgatcgactc agacgagtaa gatagtaacg 360
 gaggatccta gtcctangga tccttgtac canaagtctt tctggcgatg ggt 413

<210> 30025
 <211> 461
 <212> DNA
 <213> Glycine max

<400> 30025

tctcttgatt cttgaatctt cttgatttct tctcatgaaa cttgaaattt atcttgatct 60
 tgaacttggt gactaaatct tgaaatcatt ctttgggggtt tttgtcgtca tcttagtcat 120
 catcaaaact tcttgaatca acttgattca tcatcatgaa gcttgcttct acacttaacc 180
 cccaagacca aaaaccaact agcctgagag gctaggaaaa aagagccacc agtccctcta 240
 aaagagcccc catatccttt agttccatca aagaagaata aggagcacta cttcaagtgt 300

ttattggaga tattcaaggg gttggagata accatgccat ttggggaagc cttacagcag 360
 atgctgctct acaccaaatt catgaaggac atcttcacca agaaggggaa gtacattgac 420
 agtgaaagca ttgtggtggg aggcaactgc aatgcagtga t 461

<210> 30026
 <211> 364
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30026

agcttgccgc ttctgagnna nactactatg tttcttgngn ggnggaacaa gctacaaaag 60
 gatagagcaa gaaatgaata gccaatgggt gatacatgga cggagatgag aaagatcatg 120
 aggaagcggg atgtgccggc tagatactgc aaggacttga aattcactct ccgaaatcta 180
 acaccatgca acaaggaggt tgaggagtat ttcaaggaaa tggatgtgct gatgattcaa 240
 gcaaatattg actaagatga cgaggcaact atggctcgac ttattaatgg tttgactaat 300
 gatatactg atacttgtga gctgcatgag tatgttgaca tggatgatct gcttcacaaa 360
 gcaa 364

<210> 30027
 <211> 438
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30027

actgtgctag agagnaaaac aaatgaccaa agtgaacata gttccatttc tagggcaaaa 60
 ttgggtgttg agaagtcaaa ttttgattcg gtggaatttt acgtgtaaat ccagtttgag 120
 caagtttaga ttgatgttat ggacttgtgt gaggagagag tttgcttcaa atttacctca 180
 ttctaaattt cacttctcaa gcctagaaaa tccattaaat tgaggggttt tggacaccta 240
 gattttgtgt tgctgtggtt tgaagcttgt ctttggttta tacatgattg atacatgatt 300
 tgagacttgt aggatttgat ttgggcaaga ttggatgagg ggaagtgtga ttgtcgaaat 360
 ctgcactctg tgcagattat tgctgtgaaa ttgtgcagca taatcttgca tgagtgcata 420
 caaatgcttg tgtgtgat 438

<210> 30028
 <211> 328
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30028

 ctcacaactt tcccaaaaaa gaagtttaaa gatgtgggaa gccttcatcc accttgtggc 60
 atggtataaa agtgtgccat cttgctaata caccaccata nagatagagt ctacacctct 120
 ttgggttcta tgaagcccaa tgacaaagtc cataactaatg tctaccaag gtgcanatgg 180
 aatgggtaag ggtgtgtata gcccatgagg catcacccta gacttggctc gtaaacaagc 240
 cccactccta gtgcaatgct tatggacatc tttcttcata tggngccaat aaaacttctc 300
 tttgagtaag acaagggtct tgtctatc 328

<210> 30029
 <211> 449
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30029

 gctttgtcac aaagaagaag aagatgttca aagagattca aggcttgtaa atgattgtat 60
 aagatctatt ggaaaagtat attgaaaagc aaatcaaagc cttgctttta tagactcttc 120
 atgtctggcc aagaggacca tttagaagag ttataacttt tagaaaaact taaaaccaat 180
 ttgaaaaagt caaaaaccat ttgaagagtt acatcttttg atttatttag aaacaatcac 240
 tggtaatcga ttaccaaata agtgtaatcg attacacaaa gcttttatgt gaaaggatga 300
 gactcttcac atttgaattt gaatttcaac gttcaaaggc actggtaata gattacacaa 360
 acattgtaat caattacagc tnttttgaaa tcaattggaa cgttgtaaata tcatttgaaa 420
 aaaatagtgt gtgcatgcta tntcattat 449

<210> 30030
 <211> 446
 <212> DNA
 <213> Glycine max

 <400> 30030

ggacacttga tactaagcta cacacacggt cactgcatgt ttttgcacag aatgaagaat 60
 atttaaccaa caactttgtg caagaaatcc ctacacacac acacacacat attaataata 120
 aattgaaacc aacttaatta aaacaattta aaacattctt tttaaaatac aagcctttca 180
 aaggggaaag gctccattac cttttaacat cataataaaa cttgtacaaa taaataataa 240
 attcacttcg gtcataaca aggcgggtcta aaacttgata caatcaacat agaacctata 300
 ccctaattgc acatcctatc ctatcagagc attgtattcc cgtgtgctct agcatcaggt 360
 tcttcatagt catccaccta ttcatttgct ccactaaca ccacgttaga gatcatcaca 420
 tgatccgaac acagattata cactgt 446

<210> 30031
 <211> 331
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30031

tttgcnnga acttttattc atgaaaaagg gtctttgtta cgtgtgggtg agagatgtta 60
 gaatgagtaa tattgataga agaggaggag atatgagata ctacctgcgt actttacaac 120
 aaaggccaac atgggtaaag cttattggat atatggcggg aaacatatag cagacaacag 180
 cacgaataag gagttcagcc agaccatagc tatatagata aaaatctcaa aggaatatac 240
 caagtggat ctaccaacc catgcaaagg aaccacgtgt ccagtgatta gcctaacgaa 300
 ggtatcactc ttggataagc aacgagtgat c 331

<210> 30032
 <211> 442
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30032

ngccagctgg tagtccatgg acatgccctg aagacgaagt gttttacaat gcttcaatca 60
 tttggggagt agtgggacca cagagaatgt ttaccaagga tgggatttac cctgngatga 120
 attggttttt cctgattggc ctacttgctc ctgttccagt gtggctgctt gctcgcanat 180
 tcccaaacca taagtggatt gagctcatca atatgccctt aatcattgct ggtgggtggg 240

gcattcccacc agccagatcc gtcaactaca taacttgagg atttggtggga atcttcttca 300
 atttctacgt ttacagcaag ttcaaggcat ggtgggctag acacacttac atcctctcag 360
 ctgctttaga tgctgggtgtt gctttcatgg gtgtcattct ctattgtgcc cttcagaatt 420
 atgggtgtttt tgggtccaata tg 442

<210> 30033
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30033

ttttgcttat agattccttt gctttttttg ctagacttga agctataaga atcatgcttt 60
 cctttgctac tcataaaaat ataaagttat ttcaaagga cgttaaaagt gctttcttaa 120
 atggctttat tgaagaggaa atatatgtca aacaacctct tgggtttgaa gatcatactc 180
 ttccagacca tgctttcaaa cttaaaaaag cntngtatgg tctaaaacag gaaccacatn 240
 gctgggtgtga cagactgagt tcatttctct tagaaatggg tntattaaag tcaaagtggg 300
 tacaactctt tctaaatgag aaagtggcan agatttcatt atagttcaaa tntatgttga 360
 tgatagtatn tttgaagcta ctaatgaatc tctt 394

<210> 30034
 <211> 451
 <212> DNA
 <213> Glycine max

<400> 30034

tgaatgaata taagacacat cttcttcaat cttgggtgatt cttgactcca tctaattggaa 60
 gtgcatgtcc acttgtaatt ccaaagtgtc aaacctttca ccaacaaagg tttgaagacc 120
 atcaaacctg tccaaaatct ttgaaagaag agatgaatct tctccatcat gtccttcttc 180
 accaatatgt cgagcaccct ttttcaacca agagccatca tgctcttttt gataacaaaa 240
 ggatgcaatg actgaagcgc ctataaggaa ggatctcttg attggaacat aggggttcaga 300
 atcaagaggg atgttaaagt gttgaaggaa aagagtgact agatgaggat atggcaaagg 360
 agcattcaat cgcaatgcct tatgcctgcy atatctaaca agaagtgccc aatcaatttg 420
 tagaccttta tgataggccc acataacaat g 451

<210> 30035
 <211> 382
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30035

tttgctaata acttataaca aatcaacgnt atatccttgt caccgaaaat attttatatt 60
 tttttctcgt atccacattt actcgagatt atttcttatt ttcataatttt ctaataatttt 120
 caaagaatgt gccactcata aagtaacatt ccaaattaga gataggcatt catctacttt 180
 ctatgggaac attagtaaaa cacatganat tatntactat gtgttttaa tgtgccgttt 240
 ggcatacat gaacaagggt cttcatatca cgtaaaaagt agataaataa aagtaaacia 300
 ataagtatgg catatcccat tagtctaaaa gcaagggtta tatattcaaa ggtttatatc 360
 ccattatttc atatgttcac at 382

<210> 30036
 <211> 446
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30036

tggttcgagg tacttaccg ttgaagatcg aagaacgatg aagaacgaat gaagaacgtc 60
 gaagaacggt tgaaaccttt gcgaaattct tcacagaaaa cgttacggaa acgtttcgga 120
 agcgccctcg cttagatttt cttcacggaa acgatttttc caagcaaatt cgaaagagag 180
 agaagtgcca aaggggctga acattttctt cttcacttcc tcccctattt atagcaaaat 240
 aggggaggtg gttgccgcc agctcgccca ggcgagccag gttgcttctt ccagaagcaa 300
 cagccttctg gaggaatatt ctagagggcc caagtgggcc tgggtgctat ttccaccccc 360
 atttttacta agtacacccc cctctgctnt tttggtgatt cttntccgt anagtcacgg 420
 aaacttacga attccgtaac gatact 446

<210> 30037
 <211> 509
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30037

aaaagttggg ccttgttgan cctgagngct tttcggagat acanggcgaa gttgagcacg 60
ggacccgtgg atactacaga gccgaccttc ttgcatgcga gctttgacat tggctaggan 120
gcagatgagg catacaacga aactctgcgt atgagagtta aacaagagtg aatcataagt 180
acgcatgcc aactgagtaat gatgaaaatg agaatcgaga tgctgaagat gatgttggat 240
tggaagtatc aactgtggac ttggaacgta cagtgcgatg aatggctctt actctatgga 300
gttagtatat gtcaaggact ggacaaaggc taattttcat gaccgaaata accctgtgaa 360
acatgatgtt ctatctatat actgtttaat acgaacacag tgtgtgctat agatgtgata 420
tcgatgcaaa gatgtgctag cttctattaa tgtgacaact tcttatagct acccatgcta 480
tgataatatc atatacggat aataacgtg 509

<210> 30038
<211> 428
<212> DNA
<213> Glycine max

<400> 30038

tatgaggtcg gttatatatt cttctttca ccctatttga ggtctctttt tcctttcata 60
agagaaaaat tgtaaatttt tagtctctca tttatatata taacacaatt tcatacaaga 120
gaaattaaaa agacattgat ttatttctga aggattggaa gctaacattg tcttgtacta 180
aaactatatt tgacaccttt tattgcatga tcgctttctc taatatagaa tctaagttag 240
tgtattgggt tagtgcaggt tctgagtgat cccagaaga gagcaatcta tgatgaatac 300
ggagaagaag ggcttaaagg gcaagtgcc cctccagatg ccggtggcca tacattcttc 360
caaactggag atgtgccaac aacgttcagg ttcaatccaa gaaacgcaga tgacatcttt 420
gctgagtt 428

<210> 30039
<211> 133
<212> DNA
<213> Glycine max

<400> 30039

ttcgagcgtc tccatatatt atgcgcctta atcggaagat cgagtgaaat tgtttgacca 60
 tttgaatgct ccacagcctt ctatgggtcaa attcgagcat cttgaatttt atgcacctta 120
 atcggacctc cca 133

<210> 30040
 <211> 398
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30040

ttgaaaagtg ttgtttntca ccttctcgct acgcctttct actggcttag cgagcatccg 60
 ctaagcgcaa cactcatggg cttagtcaa ggaagactct cgacgaagat gagttgcaca 120
 ggttcgaaa ggcactgtt tcattctact aagcacaccg cttcagtcca tacgctaagc 180
 gagaaaggca tgtgctaagc caaaattcac taatgtgcgc ttagcgggtcc attattgtgc 240
 taagcgcgtg agcactatca aggctaccta tataagccat aaatcatgat ttgtgaacgg 300
 agtttgggct gtgattcaga gctttagatg gttagagatg ttatagagag aaagtctcag 360
 ttctagagag ttttgagaga tttgttggt gatgatct 398

<210> 30041
 <211> 408
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30041

agcttggttg ttacaatgac attgactggg ctagagatga agatgatcag aaaagtacta 60
 gtggatatat gtttttcatg ggaaatacaa cttcacttg gatgtcaaaa aagtagtcga 120
 tatatagtca ttcttttgac ttgtaagcca aaatacctag cagttgcttc atgcatttgt 180
 catgcaatat ggctcaagaa tttgttaaaa gagttgggca tgtcacaaga agagttacca 240
 agatctttgt cgataattaa taagtcagtc attgctctag caaagaatcc aatgttccat 300
 gatcgaagca nacatattga taccggttac cactacataa aggagagcac aacaagaaag 360
 gatgtacatg cangatatgt gaagtctcaa gaccaagtag ttgacatc 408

<210> 30042

<211> 277
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30042

gggacgaggg caacccaaca ctatggctac attgcacacg aggggaacgcc ttaacgtccc 60
 cgccaccact aggggaagaa tcatggtgac cagaacaagt tcccaaactg gagaagtagt 120
 ggaaggttgc accctccgc ggaggtgaga agtgctcca ccagcacgac acaagaccgt 180
 cacactcttt gaggaagcgg acttgctatc gaaagcccca tggcgagtct cctccganga 240
 ggaaccgccc atggagacga cccttagagt cactatc 277

<210> 30043
 <211> 480
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30043

agcccggttt atgcttcac ccnaatttg nnanngacgg ngcgngnnn aagctatatt 60
 acaannanaa ntaanatgnn tttccgaagn gggacataat tgggcaagg ttgcctgttt 120
 ttttctctct gccccggcgc cacatgccac atgcagggat ggtgggatca gtatttgaga 180
 gattactatt aggagctgaa tttgagagac atgtgccagg aaaaagaggg agaaggataa 240
 agataaacgg atagaaacct tataatgaat aaagtctaag aaatgttaca agtttgaatg 300
 tgaccgtcta atgggtatga taaaaatctc aaaagttcga atgtgattgt ctaatggcga 360
 tgatattcat attttctact aaaaaatctc atanaattta tcagaattta tttaaaaaaa 420
 cattaaaatt gaaaactttt gatatcaaga gacttttata aatataaaaa tctaattaat 480

<210> 30044
 <211> 405
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30044

agcttctcgt ttatgagaac aaatntcaat atcattcatc ataattggct cccaccaatt 60
 tttcagattc aagtttacca cagaatgaca atgaagacat actgttaatt aataacacat 120

ccaaagcggc gacatattat cttgcaacca cttaagctta aagtagacct actattgttc 180
tccttttgga gaacaacacc canatattgt tcatacatTT cctcccaatc ataataagtt 240
gcaccagtaa ttggcctatc gtccaccctt aaaccaagct gtagtggtac atcctctagt 300
gtaattatac attatccaac aagaaaatga taagtgtgcg tctctggTca acaagtgcag 360
gcactagatg atgatcaatc ttgaagtgtc ataattttgt cacat 405

<210> 30045
<211> 306
<212> DNA
<213> Glycine max

<400> 30045

agataaatag taaaaaagtg ttttaaaaca ttgagtagca caagaatttt tcacaaaatc 60
ttttaccaa gagttctact ctctggtaat cgattaccag aaggtagtaa tcgattacca 120
atagccaaca ttgtttttaa aactgattta caaagttgta atcgattacc atgagcatgt 180
aatcgattac caatatttta aagcgttaga tatcaaTgt cagaagtcac agatagtgat 240
agaacatttt caaaacagtt taaacttTgt taagcgatta cacaatactt gtaatcgaat 300
accagt 306

<210> 30046
<211> 403
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30046

agctttcata tttgaaaaaa aaatactaac ttttggctct tggaattttt cttctggctt 60
tctaagatat gaactggtaa gttgttctaa ataataaagg atatggtctc agattatttt 120
gcgcagttga aattcttgca tactatccag ttgcaatctt gttagcaagt agcaactaca 180
atgacactta tanacgatgg aaaaatctat ntgactactg ntgtgcatcc ctttctaaag 240
aattgtctgt gcctgtcttt ttgtcctgct tgtaagctgg acanaatgag gaatattctg 300
aattcattat ggatctactt cgtaagtcct taactgactn nttacttatt tctttgctgc 360
agggagatca attgcaagtg acagtaaata aattaacatc aac 403

[illegible]

aggaacggcg	gataagaata	gagagagaga	ttatgtttct	gggaagtgac	tttattctag	60
tagccgcgtg	acatgagcct	tcgcattcgc	ccagaataca	ggaacaaact	tggaatgggc	120
accaggcgca	accacgtaag	cccaattctg	ttcaaacaat	ctagttccgg	gtgttactag	180
gagcaccag	aatttttttc	tggggcacc	aaaaacatag	tggaagaata	aaacaatgag	240
taaatagacg	ttataaataa	atagtattgc	tatatataaa	actaatcccg	tgtttaagaa	300
cgctcaaga	atttcgagcg	aacctagcca	gcaaaagttg	atggaatttg	gactcaaaag	360
aatggcggtg	caattctaga	ctatagttgc	ttttggcaaa	actaacgcta	aacaatctat	420
tggtga						426

<400> 30048

agcttttttc atcttatcat tatcaaacag atggtcagac tgaacgaacc attca 55

```
<223>      unsure at all n locations
<400>      30049
```

tatcatgatc	tggaaaagta	aaacaatgtc	agttatatag	tttagataat	catgattata	60
tcagtaat	cattgcctcc	aatatcatat	aaagcattta	gtataatata	taacaatata	120
tggtaacata	tgagaggttaa	aagcttacaa	acatacgctt	acaagggtcat	ttcatatatc	180
acaaaattga	agataacatt	caatgggttca	tcatcaa	atgttgcatcata	ttaatatata	240
cattttgaat	aggcaacact	tgctctcag	tcaaaatgca	ttttcagcaa	ttggatgtta	300
tctctaaaaa	agttaagatt	gtatcagact	tagtagcaca	ttcagcagca	tgacaaaaga	360

cacaggataa ataatgcacc atangatcaa taccaaactg gaagagataa cttctcagtt 420
 ttaaaggtga acactattat aat 443

<210> 30050
 <211> 408
 <212> DNA
 <213> Glycine max

<400> 30050

agttagtaat gagtggaaact tgatttatgg tttagctcaa tttgacttaa aagaatgact 60
 tagttcaaag cttgttacaa gtttaatatc aacttttatt ttttatttga attcagtttg 120
 atttaaactt gtgagtaatt caatttagct cttttgttgg attggtttaa aactataatt 180
 attttaattc ttttatttat ttatttattt tgtgaaataa aattaattaa caaactaatt 240
 atgtcaattt actattttat atcaatttag tcatattaac acatgtaaaa ttagaggatg 300
 aaattcaggg agaaatgaca ctggctattg gcgtgaagtg ggcgaaacag ccgaaaacct 360
 gaactgggac atttttgcac cacaactagt acttaagtag tctttctg 408

<210> 30051
 <211> 404
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30051

agtctttatt tttatttctt gacnatcttt aatgagctgg tgtaacagaa aatgataaaa 60
 taatcatgta aggtccgttt tatacacaca tacatctagc gaattctaatt ttacatgtag 120
 cggtttctcg acaaaatgca gatatgcttc tgataattgt tttgggggtgc attgacaccc 180
 atcaaaatgg gacaacaatt gattggctcg tcaaccatgc gcaatcaatt tattatcaga 240
 taggttgcaa cgcattgatt ttgaatcctt gatcgttacc acacacgtgc cctatatgag 300
 caactatcgt agagaccaag aactctttcc ttgaaataag ttccctcttt cttttctgta 360
 gtgctttgct tgatcttcca ctcacggtga taaaaatctg actc 404

<210> 30052
 <211> 419
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30052

gaccgctntc agttttatatac ctagtaaatac catagatang tttatcaaac cagtatacaa 60
atggggacag aaaatgaaaa aaggtaagtg agctcaaact caaagaatgc ataatatatc 120
atttcaaaat cagaaccact gaaccaccaa acattataaa agaatttgcc agatcaaaca 180
tctagaatgg gtagacctta caagaacaat tagacgagaa gaaagtttca tgcagaatta 240
atgtacaaaa tgcaaaaaca aaagacaagc cttcaaaacc tatggtcaga gacactacgc 300
ttattggatg ttattcatga acttcagtat ttgttattag aatcatatag tttcaggagt 360
cttattaata tgctatgta tcattgttat gtagcagaat tgtacaagac ctatcatat 419

<210> 30053

<211> 367

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30053

ttactcggaa ttggtaacta catttttta gctcgaagtt ttactgaatt ttgtagacat 60
ttggaccaca attataaaaa aagaaccaag cgaattggat taaagaaaaa aactaaaaaa 120
atcacacaag ttggatgaaa aatcagtgct caggaaaata aaagtgaaaa ggaagtgtgc 180
ttgttggttt aactcaaac tttttctata attgggtgcct actttatacc actcctagtt 240
ctgaaacttc aattgaaaat aattatgaaa acaagtgcc aaaaatagagg tttcttgagt 300
ctttttttcg tttctcttta ttaagntttc tactctactc tatagccttt ctaggtttgt 360
ctttgag 367

<210> 30054

<211> 427

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30054

ctaattatat cctataaant tgctatcact tactacttac atacattcga agtacaccat 60
acaaattttt gttgtttcac tcctatttat ttatatgcat attggaaagc taattacatc 120

ctgcacatat ttgcattcaa aaagggcatt ccacactatc atacattcat ttaagaaaac 180
aattactcat actttgctag gaatttcatg ctctttatat ttacctatgt atacacacta 240
ttgcaaggtg ttttccacgc tacctctatg taaagtatca aacatggggc agcccaaatt 300
cgagcaaaaa ctctcacaag caaatcctaa ttttcatggt tttctaattc taaaacaaaa 360
ttntggattc ctagccataa gcatgtttcc ttgcattgaa gctacaagtt tgggttccta 420
agcttgg 427

<210> 30055
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30055

gtcatgcatg ttatatacat gaaattgatt agactattat tttattcgac cggnaaaata 60
tcatggacat tgaaaacatg gccacaaaag tcacctccat gaaaagttaa aagaatttaa 120
ctaattccta taactaatat attttaagag taaattaaat cataaatctg caaattaaaa 180
taaaaaactc aaaaaaagaa aacaactatt aaaaaaatac aatacataaa tatagaatta 240
aaaaaataaa ttcataaaac aaaaaaatgg cacattgaga aattgggttg cgacatattg 300
tgtagcaaaa aaaattaaag ctggacagtg agaaatcgga ttaggggcac cggattttca 360
ttggcaataa tgatttgtat cacttttgat agataatttg gaacttgtat tatttgt 417

<210> 30056
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30056

tgtatttctt gaggaataac tgtaatactt aattgggtgc acctacatgt aacaacatca 60
ctttccccgt caacaaacat gatcaaatca acaactatta tttctacaca ttaggggtgaa 120
aaagcactca atatttttgc tataacataa caaaagtgtt acaacaagag taaagtgatt 180
ccaacttcaa cttttttttt tcaattccct catttcattt ggggtgcctaa ggtatctagc 240
ctagagtcag aaactaattt ctcaagacac aaagatcgat tcaagggagc tgaactcacc 300

caacatanaa acataagaga ctcaatcact tcctaatttc catccnctaa taaaacatga 360
 tttgtggaac aagtgaagta aaataactaa nagaattgaa aagcacctgc atgttgact 419

<210> 30057
 <211> 276
 <212> DNA
 <213> Glycine max

<400> 30057

tgaacagatt tgataggagc tagggagttt accctctgat tcatataaaa aaaaaagaca 60
 tcacatcatg cttaccggtt taataggcat ggaatagaca agtctctaag ttaacacggg 120
 aagtgcgagg catgaagcgc caatcgcgag ggacaagtaa cccataacca tgaactgaac 180
 atgaagaagc ctccctaagtt aaaaataaag cgcaagacgg tgggataaat tgggattagt 240
 acagcgttgt aggccaaaat aaaacgaaaa gcggaa 276

<210> 30058
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30058

ctgctactta tatatcgaga cgatcaaat tgaacaacgg aagctctcgt gaaattaaaa 60
 tggtcataag ttttaactcg gatgtccgat tcaggagctt cacatatcga gatgcacgaa 120
 attgaacaat ggaagctcta gagaaattct aatggtcata aattttcaca cggaggctct 180
 attcaggcgc ttaatatatc cagacgctcg aaattgaaca atggaagctc tcgagatatt 240
 caaatggtca taacttttca ctcggatgtc cgattcaggt gtatcacata tccagacgct 300
 cggaatngat tagcggaagc tctagagaaa ttcacatggt cataactttt cacacggatg 360
 tcctattcaa gcgcttaata tatcgagacg ctcgaaattt gacaac 406

<210> 30059
 <211> 389
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30059

acaagtttct tcacaaataa ctatcatgtt gtctgtttac tagcaagact acccatcata 60
tctcccaaaa ccccataccc acgaaattta agagagaaag aagtccaccc aaacctgaat 120
tttcgaagtc ccactcgtag ccacgcactt cagacccccg aaaatgccct cttttcgga 180
tttggggcag aaatgatgga caaagggtga agctttgctt ggagcttcaa tggagaatga 240
agaagaagaa aatggcaacg tgagggagag agagagctgt ctganaaagt gtggtgctga 300
gtgaagagag agaacagctc tctgggttta aataaaaggg tttctctttt ctattatttt 360
attaagcatt gcacatgtct catttgagt 389

<210> 30060
<211> 343
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30060

tagctttgtt tactttattn ttggtgaagag gctttacata ttatatacta agatatctgg 60
ttgaatcttc catttcttat ctatggtctt gaacacttac gaagtctacc agtggtatgc 120
taatcatgtg tatgcaattg actagatgaa actagaagga atttggtaga ttggcctaaa 180
cgatttaaca ttatttgtgg cattgctcga ggacttcttt atctncatga agattctaga 240
atgaggattg tacatagaga tctgaaaacc tagcacattt tactagatga aaatttcaat 300
ccanaaatat cagactttgg cttagcacga gcattcttgg gag 343

<210> 30061
<211> 419
<212> DNA
<213> Glycine max
<400> 30061

aaagtctcac gattgtcacg tgctcatgca ttatttggtta gtcgtggcta tacgagacat 60
cttgcgaaac aaagtcaggt tagcgataac tcgcttggtc tttttcttcc atgctatatg 120
tagcaaagtc cttgatctag tcaagtttga tgagttggaa aatgaggccg caattatact 180
gtgccagttg gagatgtatt tcccccccg ctttcttga catcatgatt cacttgatta 240
tgcactctggc cagagaaatc aaatgttgtg gtcctgttta tctacggtgg atgtaccag 300

ttgagcgata catgaagatc ttacaagggt atacaaagaa tctatatcgt ccagaagcat 360
 ctattgttga gaggtacatt gcagaagaag ccacttgaat tttgtcataa tacttacag 419

<210> 30062
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30062

agcttatatt cctattgtcc tatagaggca tttcattcct tctggagagt ccactagctc 60
 tatgatagtt gtcacacctc tgaacatatt cataagcatc cttgtgcaag gtaagccaat 120
 aaaaactaga ttgaaggacc ttggcagtag tctctctccc atcgtaattg ccttcacaat 180
 gtgaactatg gcaatgccac aatatgcttc ttgctcccc ctaagttaca catcttctca 240
 agagattatc tgctccaatt ttataaagat tgggatcccc ccacacaaaa tatttagtgt 300
 ccttgacaaa cttctttttt tggaccaagt gagatcatca ngaaatgcac caattgcttt 360
 gacactagtc atctcagcag accatggcct tctacaatg 399

<210> 30063
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30063

tggtcttcgc cagtgaaggg atcaatgtgg ttcttattta tgcaaatang atcatcctac 60
 tatgacgact gagaaaactg gggcaaataa agagggtgag gatgaggcac aaacccatgc 120
 tgtgactacc attcctgtac ggccaagttt cccaccaacc caacaatatc tttactcagc 180
 caataacaaa ctttctcctt acccaccacc cagttatcca caaaggatcat ccctaaatat 240
 accacaaagt atgtctaccg cacttccaat gacgaacacc acctttatca caaaccagaa 300
 tacaccaacc aagaagcgaa ctttgcagcg agaaagcctg gaggaatcac cccaattcca 360
 gtgtcctatg ctgacttgct cccatatcta cttgataatt caatggtagc cataacccta 420
 tccaacgtta tca 433

<210> 30064

<211> 251
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30064

atggttaata atctgcataa aggtgtgaat gtgtgtggta tgtttggttt agttcaattc 60
 ttggtgcatg gatngagatc ccacattgac tatagatatg gctaaagtag aaattataaa 120
 ggctggggaa atcctcacct catgaagcta gctttggagt ttgagtcaag cttatctcan 180
 attcaagatg gtatcagagc ctatcataaa ttcgatattg ggccaccctc aactgtccaa 240
 aaatctacat g 251

<210> 30065
 <211> 431
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30065

aacctcgtgt aggcgcttca cgcncgccat gcgctttttt cgcgtncctt gcgcgcgcga 60
 gctcgagcgc ttcctccatc ggatctggcg cggccgggtg cagatctggt ttaactgtca 120
 ccggattcac caccgcataa cggcgccggg aagttgcgaa atgcaaagt aaatgcgaat 180
 gcgggattta aaaggagtga gagggaggtt cagtggagt tagtgagaga gagagattgt 240
 gaatttagtt agcggaggcg tcaaaatgag gggagaggtt gcggtacgag agagaaagag 300
 aatgatgacg cggaaatgaa atgaaatgga aaatgatttg aaaactatgt tttatgctat 360
 ggggctgccg tgcttgaaaa ttaaccacta ttagtattga gtgatagttt gttgggttgg 420
 gttgctttta a 431

<210> 30066
 <211> 478
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30066

cgggccggtt cctatgccat gaatatctag gcaattcagc tcgtaccgg ggatctctaa 60
 agtcaacctg anggcttgca gcttttttac attattgtga gagtggttga cagaaatcaa 120

aagcttattt tggaggcccc ttttttaaaa aataaaacat ttaaaattga aaatgatggg 180
ataaaataaa aatgggttgg cactggaagt aaacgtggag aatgggtatt ggattactga 240
tttgccctt taaattttta aaatctgaat cagcttaact caagaaaaat ggtgggtgggt 300
ttgcctcaga tcaagccttc tagtgaagat gtgatggttg ttacagtgtg agcatcaaga 360
gcactttcaa acaaatgacc aatcaggcaa aagagaactt gaatgattac tctgtgtgtg 420
tgccctatgc gactgatctc tgggtgaaca gatctcatat ctttatgatg attgacan 478

<210> 30067
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30067

ctacaagaga agatacaaat tctaagaaat tctttaattg tgtgatgaga atatgttata 60
ccatgaactt attttctaaa atttttgcaa ttggtataat tgaagccttg cttgattcct 120
gtttttcttt ttcttttctc atttatgttt tgccaaggca tttttttctc atttattatt 180
ttctgtcatt gcattgagat cgtaggcata gattcaatct tttcctctaa gtacacacat 240
tctaaaacag atttttaatc aactgatgag aacagggttat accatgcac attattgtta 300
gaattcatgg ttcagttcaa gatagaagag agaattgaga aaaaaaaaat tatatgtata 360
ttaatattct taaatgagta cccaacaagt atataccana gattccctat gagatgatga 420
ttgaaaagaa gtctag 436

<210> 30068
<211> 117
<212> DNA
<213> Glycine max

<400> 30068

tgattctgat ataataatga aaacttgaat agtgatgatg attcttgtac caccacagct 60
tttactagga aatcctccaa agtcatttcc tcaagtgttt gcttgctttt gagcagt 117

<210> 30069
<211> 444
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30069

aatagattta gttagtcact ctacttgcaa aagagagatt ttctagcttt ctaggaactt 60
agacgttgat gaactttgtt taagataaat tgaaaagtat tcctagaagc tatcttatga 120
aagatagaca ctccaaggta ctttccaaga tccttagtcc aagcaatacc catttctcca 180
cttagttgat ccttgacttg agtctccaca tttttggaaa agaacattca agatttctcc 240
aagctaattt tctgcttaga actcttgcaa aataaattca aaatattctt gatagaatgg 300
acctgctcca ctaaagcctt cataaataaa ataaggctgt atgcaaaggc taagtgagat 360
ataagtggac catgtctaaa aagacgaata gggcaccaca ctctntggtc cacaacaaca 420
gagatcaatt gaaacatatg ttca 444

<210> 30070

<211> 407

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30070

agcttttgatt tacacagagg gtttcagggg caaagcttcg atttacetat tgaatttcat 60
ccaatttttt gacaagtcac tgttacttcc atgaatattg atattgtatc atgcttaatt 120
atatgcattt gattattctg atcattgtgt gttgtgtgat tatttcttcc atgcaggtag 180
atgattccta tttgttgtga gaggtaaag atgggcagca tcaccaactg aggtgagttt 240
atatttcctt ttttttgtct ttatctttgt tagttcgtaa tatagttttt attttatatg 300
tttgagttct acatgtgtaa aaaatagaaa tagacaggtc tgggtgattgc ttangaattc 360
cttggtgttt catggcattc ctntgaacc tcanaagggtg cttatga 407

<210> 30071

<211> 426

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30071

gctgcacat tgacagatnt acttagtaaa gaagcattct tgttgtctcc agaggcagag 60

acaacatttg ttcaattgca gaaagtcatg acttcagctc cagtgttagc tcttcctaata 120
 ttccagctgc ccttcattct ggaaactaat gcttccgaca ctggtattgg agtagtatta 180
 catcagaatg gccatccaat agcatttttt tccaagaaac ttgcacctag agtgcaaaag 240
 aaatctgact aatttagaga gatgttagca attgttgaag ctatagctaa gttcagacac 300
 tacttgctgg gacacaaatt tattatcaaa actgatcaca attagtcaga tgatgatgtt 360
 gatggatgga acaaccgcta cagacacctg aacaacaaca gtggttacac aggttttttg 420
 gatatg 426

<210> 30072
 <211> 382
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30072

agtctttttg aaaagattcc taaagaagct agagcttagc tacacacacc tctctaatag 60
 ctaagctcat ctnccttgaga tgagaagctg gaacttagct acacaccccc tataatagct 120
 aagctcacc ccatgacaaa atacatgaaa atacaaagaa nagtccctac tacaaagact 180
 actcaaaatg cctcgaaata caaggctaaa accctatact actggaatgg ccaaaatata 240
 aggccataac gaaggaaaaa tacctattct aatatttaca aagataagcg ggctcact 300
 tagcccaggg gctcanaatc taccctaagg ctcatgagaa ccctanggcc ttccttgga 360
 tctctggccc aatctacctg ga 382

<210> 30073
 <211> 446
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30073

gggctganaa tatataacaa caccaaggat ctacttttat ctctcctctn tcgttttttag 60
 ttgtaggctt ctcttcttct tttagacact cttagccaga agtagcaaga aaaaaatatt 120
 tgttttgtaa tcaaagtttt gattagtggg tgtggaagta atgctttcca agattatttt 180
 gatgatgcc aaaaactcaag tcaagaatca agagtcaagc aagtttcaag aatcaaagag 240

tcgttcaatc aaagcaagtt tcaagaatca tagagtcggt caatcaagat tcaagattca 300
agattgaagt aaagaatcaa gagaagactc aattaagata agtattaaaa gagtttttca 360
aaatattgaa tagcacaatt ttgttcaaga gaatctttca aagaacaatc ttttaciaag 420
agttgtactc tctgataatc gattac 446

<210> 30074
<211> 387
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30074

agcttgtatc attngactta tatgctctag cttgagcgag tgttggaatt agtagttaaa 60
tgcctttcag atacaggtac actattcttg tacctagaat ataccgtgca tgtactaagg 120
aactaaaatc atcattaatc tccttataaa caaaaacag tgtatataaa tactagatca 180
gctgaataac tcattcaagc tattagaaaa gtcattctcg taatccctaa naattcctca 240
tgtaataaca ttcaaccttc caacaaatgc atatggagga tttcacattc tcaattcatt 300
gatcttcatt caagtgtac taaatctcaa aatatantaa atntatgtct ttggtgcatc 360
tatcattgct taatcanggg gttatgc 387

<210> 30075
<211> 447
<212> DNA
<213> Glycine max

<400> 30075

caagatgagc ggtcatcctt gcagcccata attcggttctt ctcaccatca aaaattgggtg 60
gtgaacctgt tgtgtatgat gtttctccct ccattgattt ctctcaactc acagatccct 120
taaagataag agctctgata ccaatttggtt gtttttggtt aataacggta agcagaaata 180
ttaaagaatg aaaggagcg taaagaaaga aattgagact acaaaggctt gttttattct 240
gatatgaagc aacgtattta aaaacatgaa aggatagtaa cggctaacia aaagataaca 300
ccactaacag atcatgccta gaaaatagga tcaaaactaa tttatcctat cagtcaacat 360
gactgttatt tttccttaaa aatagcacia gaatcttato tactatagtt tgtagacag 420

tttcaacagt cacatcttaa taaatta

447

<210> 30076
<211> 326
<212> DNA
<213> Glycine max

<400> 30076

gtggtaatca gagcacaaga gcttcaagta ggtgcttctt aaaccctcat taattttttt 60
ttccttacct tctcttccat tgttggttct tcatTTTTct ccacgtatct cctcacatgt 120
cttggttctaa atgttggttaa catgaatctt tatagtttcc accgattaaa cttgctatag 180
aaactaaatt tgattttcta tgggtcatat ctcttggtct tggctctgaa ccatgaattg 240
tggtgagttt atgttccttt gagctttgtc ttgttatttt tgggtggctga aacctaaacc 300
ataaaattct tacaaaaata tttaaag 326

<210> 30077
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30077

ttagaggana ctcaatcacc taaatcatct gcagcatttc tgtggttaaga gaatccgaag 60
taggtgcatc agaagctcat atcatggctg aagatcaacc acgaacggtt actcttgaag 120
attattctag ctcgatcgtg ccacaattct tcacaagcat tgcgcggccg gaagttcagg 180
ctcacgtcat cacatatcct caatccttga ttcagctgat tcaaggagat ttatttcatg 240
gattgccaaa tgaagaccct tacacacact tggctactta tattgaaatc tgcaacacag 300
taaagattgt cgggtgtgcca gaagatgcag tgaagctcag tttgttctca ttttctttgg 360
ctggagaagc taagaggtgg ctacactcat ttaagggana caatttgaag acttgngatg 420
aggttgtaga gaag 434

<210> 30078
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30078

tagctgctaa actaaaatca attgagggaa cctccccag tattccatt gaaaaacctt 60
tatatcaacc tttcaaagtt agtgaaaagg ctaaacgaaa aaattaggga cctataaaaa 120
actaattctt aattgaaggg cgacgtgata accatagtga aatacttaac aagattgata 180
gtttacttaa aggcatttca gatactcccc aagcctcgga aaatactttc aaaatggtaa 240
caagaagtac cctccaatta attaatgggt ataatgaaga tagtgaccac agctcagaat 300
acacacactg agataggatc agtgtcagaa aagaatatan atccantaaa ttccaacacc 360
tgagaacacc cctccaaata tattatcaac n 391

<210> 30079

<211> 436

<212> DNA

<213> Glycine max

<400> 30079

atccaagtaa ttcttgtggg tgaagctcct tcttcttgtt ctattcccta gtggatggtg 60
cgteccctct cctcttctcc tttgccttcc gctgcatctc catggtgaaa aatcaccatt 120
gaaggacctc attggagctc aaagatccag cctccataga atcttcacaa gcaagtttcc 180
atcaagtggg aatcagagca caagagctac aagtaggtgc tccttatacc tccattaatt 240
tttttgcttt accttctctt ccattgttgt ttctccatgt atctctcac atgtcttgtg 300
ataaatgttt ttaacatgat tcttttagagt ttccaccgat taaacttgct atataagcta 360
gatttgattt tctatggttc acatttcttg ttcttggctc tgaaccatga attgtgatga 420
gtataagttc ctttga 436

<210> 30080

<211> 346

<212> DNA

<213> Glycine max

<400> 30080

tatcttgtca ttgtctacgc cgaagacgaa ggatgacgat gtacctttct ggaaggatgg 60
tgaacaacga cccctaagaa gataaagctt gctgggaggt tgctgctgac acgggacacc 120
atagacttca ccgaagaaaa cgttctttga aactgaagaa gaagagaatg ttgttttatt 180

agatatatta acttttattt tatgaatgaa ggggtattcta tgaagctcat tctattgctg 240
 ggcgcacccat caattctggt ggggtgcaccc agcaacagcc aggtgaattt cgcgcctatg 300
 ttggcctcct ttcccttaac caatgagtgg tctccaatt gagcat 346

<210> 30081
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30081

tcctagcggg ttctaattat atggacctat aaatctatat atgctgacaa tagacgagaa 60
 gttcgtggat ctctctcngg gggagtaggt gtccgccatc gctttggcct tggctagctc 120
 ttcacaaatg gattcctttg catcttgga gatgaatggc aatgtaattg agacaggaag 180
 agagagagga gacgccactt cagggagaag atgagtctag aagaagctca ccacccatagg 240
 aggccatgga taagagcttg gaggaagaaa gagatgaatg aagggagaag gagagaagag 300
 cacganattt tgtgctctaa atgagctctg aaatctgaag tttaatatc agatgatcaa 360
 agttcaaaaa aatgcacaca tatgacctct atntataccc taagtgtcac accaaattgg 420
 a 421

<210> 30082
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30082

tgcttatatt aacaaaattg ccttaatcat ttccaaatat gcatgtgaat taagatgcat 60
 caacaagaat caagccaagg ctattgtgca agcaatcaat ggggcaaac acaccaaattg 120
 attataatga tggatggctc acattctcac aaaggtaaaa tcatcacttt caaattgagc 180
 tttcaaaaact atcatgacat gtagagaaga atcaatgatt tcaagtcaca aaatgtcaag 240
 aacttttatt ttcaaaaaca ttaccattt cttgaacata tcctataatt caaagaaaaa 300
 catgcaaagt cgtacgtgca cacaaaatng acccanaata ttaaactgaa gatccgacga 360
 aactaacaac atttacagag ttacacaact aacanattaa caaaaccaac 410

<210> 30083
 <211> 427
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30083

nttgaaatca aacttttcca ctggtaatcg attatattga ttatggtaat cgattactag 60
 agaataaaaa ctctggtaac ttagaaaatt ttgagaaaaa cttttttgaa aaacaaaatt 120
 gggctatgtt tgttttttga aaaatctttt caatacttcc cttgtgaagt attcttgatt 180
 tcttctcttg aatcttgaat tcattcttctc ttgaatcttc ttgatttaat cttgatcttg 240
 aacttgttga ctcaatcttg aaatcattct cttgggcttt ttgtcatcat caaaactact 300
 tgaatcaact tgattcatca tcatgaagct tgcttctaca ccaaccacaa agtcaattac 360
 agactaagcc ttgtcttgga tatttagatg cagaaccatc ctaattagaa tccactgcag 420
 aacaata 427

<210> 30084
 <211> 276
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30084

gatgccactc tacttcaaatt tcttgaagga tatggtaacc agggaaacata agtatattca 60
 ccaagaaaac attataatgg aaggaaattg gagtcttggg attcaaaaaga acctttcacc 120
 ccaacctaaa gaccttggga gtataactat tcctttgtca attggagaag tcactatggg 180
 aaaagctctt attgacctgn gagccagtat aaatttaatg gtgctctcca tgtgtaaaaan 240
 ggtgggaagc gtagagatca tgcccactaa aatgac 276

<210> 30085
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30085

eggagcncna tgaattgagt tttcgttatg cttctctacc ttcgagtttg gagccatgcg 60

tagtgattgc ttagtgcaat tctccattct caaccctttt ttcggagccc catgaattgc 120
 gttttcgttc atgtgtcctc caccttcgag tttggagcta tgcgtagtga ttgcttagtg 180
 caattctcca ttctccaccc tttttcggag cccatgaatt tcgttttcgt tcatgtgtcc 240
 tccaccttcg tgtttggggc catgtgtagt gattgcttag tgcaattctc cattctcaac 300
 ctttttcgga gccccatgaa tttcgttttc gttcatgtgt cctccaccat cgagtttggga 360
 gctatgcgta gtgatggcct agtgtaattc tccattctca acctttttcg gagctccatg 420
 aattgcgttt tc 432

<210> 30086
 <211> 369
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30086

agctttaagg ttttangacc ttaaattctc ttaaggtgcg gatgtggagc ccaactgaaag 60
 tgagaacacg tagccctcta aagtcggggg cgggtcacc tttgaaagac gaaggcgctcc 120
 agccctctaa aagcgagggt gtgtagccca ctaaaggaga ggggtgtgcag ccctctgaag 180
 gcgaggacgt gcaaccctct aaaggtgaag acgtgtagtc ctctgaagggt gagggcggtgt 240
 ggccctctga aggcaaggac atgtaatcct ctgaaagcga gggcggtgcag ccctctaaag 300
 gagaggggtgt gcagcctcat gaaggcgaag atgtgcagcc ctctaaagggt gaggacgtgt 360
 agtctctctg 369

<210> 30087
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30087

tatgcccag tcatctatcc ctatgagatg ttgttgattt attggcgatc agaattgcc 60
 ttgcttggat tacgngttg aaccaagctc atgcttttac aaaaagggtc atcaagtcaa 120
 gttgaaatat ggaagtaacc gtcttgcaaa attggggcaa aagatgaatc gagtcacatc 180
 actgcttcgt ttactgccaa acatatttag gattgtttat gtccttggtta cttccagttt 240

[illegible]

<400> 30088

<210>	30089
<211>	431
<212>	DNA
<213>	Glycine max

```
<223>      unsure at all n locations
<400>      30089
```

12549

ttcttctttt g

431

<210> 30090
<211> 381
<212> DNA
<213> Glycine max

<400> 30090

tttttttctt aggggaggcc cttatggtct ccaatggtgg catcttcgaa tcctgcctgc 60
tcttggcatt aatactggca cccaatccac catcatcaga ccaatctctt tcgctgtctg 120
acccaaaaac actctcactc tgactaaaac tacttggact tgctgggatt ccgggtttct 180
gcttgctact tgagcctggt tctaaagcat tttccaattt atcacgaagg tttaaaaact 240
tgtgcttctt ccttttcgcg cgcaattcgc ttagtacggt ttcctttgaa gccgtcttct 300
agcatgtcta gccgaggaac gaaccttgac agaagaagga tgatgagatg aatctgcagt 360
agtagtatta gtagtagtag t 381

<210> 30091
<211> 445
<212> DNA
<213> Glycine max

<400> 30091

acgatatata ttctatcttt cttccttcct tatagacctt gtatatatgc tcataaacag 60
tgtacagggg tattttaagg ctactccact cgatcgagtt tatattcaca ccagccattt 120
ctattttcga tatcttctca gcacttcaga ataacaactt gcattttatt ttcttttgct 180
tttgaggcta aatcaggcct acggatatgc ttacagcgtg tagttctgcc atctatctat 240
cacaagagaa agagaagcgt tacactaagc tagtattaag tatttaattt aataataccg 300
tgaagggatt ccgaatttat tcattgaaag taaaacatac aaaacgggga cgatcacata 360
cagaggacca gttgaaagat gtttctagtc tagctctcaa cagaccaca ctggaaagat 420
gcttggtggt tgctggcatc tttcg 445

<210> 30092
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30092

agcttgtant cattgtctcc acgccacatc tgtaaataaa atttcccacc ctaaccaagg 60
agattgtgac catcaaggct gatcaaaaagc aagcacaata atgctatgct gagagcctga 120
aggtaacacc ctatcctccc actagggagc ttgccaagcc tcaccctaca gcggttgaag 180
gtactcaagt catgaacaaa gggcttccaa tccgagcctt cattgtttac caaacaagcc 240
tggacgatga atttgatata gatttgtggg agaacacttc tgacagaggc caaaagccca 300
tcgaagagct tgtcaagctg catgtaagga cctcactagc cttgaacaca agcacatcat 360
tgatgtccta tacaagaaca tggacctgtt ctcttgcagc catctgacat gccgag 416

<210> 30093
<211> 417
<212> DNA
<213> Glycine max

<400> 30093
tccactatgg cgtagcccat agaattctatt ttaccacagc tatcaaggaa tgtagatcc 60
attttcaatt cagttcccaa caaatcctca tcattctagt tgttgtacca atgacgacga 120
caacaacaac tattggagca tggaggatat ctgggtcaatg caattagcca attactgaac 180
gggaattaaa cctataaaca taaatataaa taatatatat aaacctaagt gtctaagtcc 240
cataaattaa gctgtagtct ctggcttaaa acatgttagg tttgtttata caagtagttg 300
gatgtttgga gtacttcggc cttttgcgta ccatcaatat ttaagaacta agttagttat 360
gctccgtaac ttatgggctc ttaataaact atatctgcac aaaattatat atatatc 417

<210> 30094
<211> 375
<212> DNA
<213> Glycine max

<400> 30094
agcttttctt atctctctgc gttgcttctt ctagagggtg cttcaatttc taaatccaat 60
ggaaccaatt cacctgcaga agatctacgc atacaaacac taacaggaac agcagttaac 120
caattcaaga agaaaataaa ttctgaacta aacaaatatt aacaaaaaat aattaataaa 180
tcaaagaata atgaattaat gccttcaaac tgaactcaac tttccaaatg gaaaaagttc 240

cccggcaacg gtgccaaaat acttgatggt cgcccctaag aatactactt atttgtgtgg 300
 ggcgagaatc taccggcgag tgcacggat cgtcaagtaa ataattaaaa cgaaataagc 360
 cgaatatcga acaca 375

<210> 30095
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30095

gagagtgcct tgaataagaa gcctagtgtt tttctctaag ggaggaagac aatgagagag 60
 agggatgggg acgtgcgaat tgaaggagat tanggagaaa agttaactt tgaagtttgt 120
 ctcacatggt tctcattcat caaaattatg gcaagtgtta cacatgtttc tatttatagc 180
 ctagcacagg ggaaacttcc taacttcctt gagaagcaag gaaggtagct tccttgggaa 240
 gctagaggaa gatagcttcc tagagaaact aaaggagggc tacttacacc catccaatag 300
 ctaagctcac ccccatgcc aatacatga aaatacaatg ggaagcttcc ttgagaagca 360
 acgaaggtag ctttcttggg aagcaacgaa gaaagctcta gaggagggga aggacta 417

<210> 30096
 <211> 404
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30096

agcttgtctc atagagggtcc aggaaggata aggcggccga agggactagt tccgctcctg 60
 agtatgacag tcaccgcttt aggagcgctg tacaccagca gcgcttcgag gccatcaagg 120
 gatggtcgtt tctccgggag cgacgcgtcc agctcagga caacgagtat actgatttcc 180
 aggaggagat agggcgccgg cgggtggacat cactgggttac ccccatggcc aagttcgatc 240
 cagaaatagt ccttgagttt tatgccaatg cttggccaac agaggagggc gtgcgtgaca 300
 tgangtcctg ggtaaggggt cagtggatcc cgtttgatgt tgacgctatc ggccagctcc 360
 tgggatatcc attggtgttg gaagagggcc aggagtgtga gtat 404

<210> 30097
 <211> 423
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30097

cttggttttaa acatgattgg tacatgattt gngacttgta tgattcaatt tgggcaaaat 60
 tggatgaatg caagagtagt tttcgaaatc tgcactttat gcagaatttt gctgttgaaa 120
 tgtgcaacag aattttgtat aagtgcagaa aaatgcttgt gtatggctgg ttgtgaaaag 180
 ggtagtacat atcgggttct gaacatttgc tagcagatcc caacgggtcaa aatgtagact 240
 tatgtactag agactgccag taaaattttc gagtcgatcc aacggttaac gaattggaac 300
 gaaggaaacg ttactggggt atttgtatgt gaaaagctgt gattntgagt tgtgttttgg 360
 gcagagtttt cttgcctttg cctgttttgc ttggttttgt gagtccatga tgattggatg 420
 tgg 423

<210> 30098
 <211> 377
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30098

ttagtaatga cccactaacc tagaattaaa ataacttaat gccattaacc ctaggaatta 60
 aaaaaaactt aatggctgaa tgtaactgaa attgtggcaa ccaaaagtca cccccaatag 120
 ccaacaagtc agccaccatt tgggtctocca aaaggctgat gcctangttg ccaattgggc 180
 ccttattaca acttgaacta aacctaacta aagccctttt agttgattaa cccaaaacat 240
 atttttggtc agccaacttt acaaggattg ggccattatt tagacagact aaacactcta 300
 aaattgaaac aaagtgggtg catttaatcc tccttcattt gggccatgat acaactcaca 360
 accttttggga cttttctc 377

<210> 30099
 <211> 385
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 30099

gaataggaag tgttatgggc gaaacttcct gcngttattg ttgaccacag agtggtacct 60

ggagatatgt cgcaggggtc acgacacctt gaggacgtca ggtgggggtgc tattgcccaa 120

aaccaagctt gaccaatccc gacccaaccc gggcatagtc ggtcagtgag aacctgcat 180

gtacctaatc aggcgagctc ctgccagtca acagataata ggaaaactag accacaaagc 240

aaggaggctt gtgggtggctg gccagctgtg aattttgtgt aatatgtgga ttgaggcctc 300

tggtaatcaa ttactaaggg tgggtaatcg attacaacgc ttattattga agacaggagg 360

ctaagatggt ctctggtaat cgatt 385

<210> 30100

<211> 409

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30100

agcttttcat aaaagccaaa ctaagcaata aagtacctcg tatttttccc tctgcctccg 60

gttgctgtgc aatcccttct acagcttgcc ctgcagcagt gccttgacca accccaggtc 120

caatagaagc aagccctacg gccaaaccag cagcaataac agaagcagca gaaataattg 180

gattcatgat aatttcctcg taacctaaat ataaaataaa gaaatagtta atgatataat 240

caaccaataa attatgactt aattnttcaa ttatcaagat ttattcggtt taaagtaatt 300

aataagaatt ccgaattgaa aataataata gttattgaac tctacgaatt acttcgagat 360

ttattttttc gtctctacct acatacatna gttttttttg tgaatatgt 409

<210> 30101

<211> 410

<212> DNA

<213> Glycine max

<400> 30101

agcttttattt agctaaaatt gaccttaaaa taaggggaaa aatcttttga tatatgaaat 60

aaacatgtca cttccaaagc cattcaagtt acataattaa ctttttttca aataataatg 120

ataataaata tctaataatc atcaatgatt catgagcact taatatcaca tccaatctaa 180

aatcttagca cactaaatct tcatcgatga tagtgaacta tacaaataaa ctataagtga 240

cattaattct tgggtgtttt ttctaagaat gtattttgtg aaaattaaat ttaacttttc 300
 tcttcacaaa ctcaacgtgt gtatcaaag atacacctat tagattgcac tgtaccagac 360
 tagtttaatt ataaaaaaaa aaatcctcat ttttctttt gtttggttc 410

<210> 30102
 <211> 495
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30102

cgaccggcgn nttatacttt cgcttgactt acgtgatact tagacacccc accttgctgc 60
 gatagagagt cttactgttt attgcatgcc tatatagctc tacaacataa ttaagagtcg 120
 tactaaaaaa aaggtaaaat ctattaagag agtcttactg tttatgtcat aatttataac 180
 ttttattacg aattgggata tatatatata tatatatata tatatatata tatatatata 240
 tatatatata tatatatata tattttgcat gagtcataac ctcataagggg ctgattttat 300
 tattgggatt atcattctat tttattttga aaattgtctc tttttctacc tcgcgcttaa 360
 gagaatttct tatactatct acttctcttt acagcactta ttgcctcctt ttctatcata 420
 actcttgata ataccactgc atcgctgtct aataatatac catgtctgtc atatagccct 480
 tattctctcg agacg 495

<210> 30103
 <211> 295
 <212> DNA
 <213> Glycine max

<400> 30103

tttcttattc gagactcgcg cctccgggta aggggccaat aatttccttt tatcaatttc 60
 tggggcacccc ccctgatat tggaaggcga ccaactgtgc cagcacaacc agaaaggga 120
 gctatcacgc agctactatg cataccgggg taagatttca cccatgccgc tgcaaagaga 180
 cgagtgtgga tcatgtgtac cgacatgacc ccttttacac agatataaat gatgttgcta 240
 cttagcaaca ttcttgccag cgaccgcaat accgatctcc cctgcccga gtatc 295

<210> 30104

<400> 30106

gattccatgg atttaaaatg ccagttaaag gcttgctttt atagactctt catgtctggt 60

cagaaaacca ttaaaagaag tataactttt aaaaaaacct tgaaaccatt ggaatagtta 120

catcttttga tttttattca aaacttatca ctggtaatca attaccaaatt cattgttaatt 180

gattacacaa agcatttttg tgaaaggatg tgactcttca cattttgaat taatttcaac 240

gttcaaacac actgggtatc gattaccana tcattgtaat ngattacacc 290

<210> 30107

<211> 429

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30107

acgtttgcag atgctctcaa aatcaatccc aatgtattga gctgaatgta caaaaggatg 60

aggggtttgt gacaaggaat gggatgagtg ctacggaaat ggttgctggt ggttggtgt 120

gtggaaagga tggcaaggga aggggtggtt caacggatgg tatgggttgc aatggttgga 180

gctataatgg cagccacaga agctccattt agcgagtggg gtggatgggt gtagtggaag 240

ttggtatggt agccatggaa gacaaccatg agctctcaat gaaagcacca aatgctacaa 300

acgcaggaac aatggggaag aataacctag cttcgagggc taggaacctc cataagagag 360

aaaaataagg aaggaaaact tgtatctatt ttntgctgtc ctctattgca tactaagcat 420

cccttatat 429

<210> 30108

<211> 360

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30108

agcccttttc cttaactttc tagggactac tcacatgaat ttggactttt ggagtggctt 60

atagaccccc cacaaagaaa atagggaaag gtaacataaa atcccaaaat tagccacaat 120

tatcaattaa acccaaatat ttgcctaaga acaaaatgaa gtaagggtgag aaaataagag 180

ccaaaaagag gtgaaatatg ctaaggagaa tagaaaaata ttaaactaag aatgctcaat 240

caaatttccc cacactttat cttttgcact cctgggcaaa actaagagaa agactaagaa 300
 aaagaaatca aactaaaggt aaaccacaac taanagaaag gaatgaacaa gacacacata 360

<210> 30109
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30109

tcgccattac atgggctgat ccactttgtc taatatcttc ctcattaataa tttgtaactt 60
 ttcttccttc gtgcacaata gcatgttcta gtcctagtct agaagctatg tgtttcactc 120
 ccatgacgtc atcctctgag accagtttca tctatatatt tgcattcattt ttgacatttt 180
 ccaaagctaa ttttatggat tgaagactag tagtgcattt gagacctatc aatcctaaca 240
 agatcagttc ttcttgctga agttgttctc catctctttg tgtataagca aaagcaattg 300
 gcttaaggtc agcatcccc atttggtgaa tcacctggcc aactttgaac ttcttatttt 360
 ccatggcatg tgtggcgttt aaacaaatta gttttacaca tganattaan aataaaaaag 420
 gaaccttcga gt 432

<210> 30110
 <211> 420
 <212> DNA
 <213> Glycine max

<400> 30110

aatctacaga gatattcttc tatacgactc ctaggtcctg aataagttgc agcaacccccg 60
 cgggagacta cacctggtac ttgtcatgct gatcaacgtc acattggcgg atgtactcca 120
 ttataatagc tctcattctc ttccaaaaga agctggatga caaccgctta tacgtcttcg 180
 tgaattccga atggccacct gcaaccgtgt cgtgaaattc ggccatcaca gtaggttacg 240
 caacatcaga cagcctttga agaataggac accatcgtgc agagtatagt gtggcagaga 300
 atcagaatca gattgcaatt gcgagctcaa cttgaccaac tcaaggatcat tatgcacttc 360
 gtgcttgata gcaggggaatt ccaccaata aggactcatg cagatgggtgc tgaactcact 420

<210> 30111
 <211> 416

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30111

 agcttggttat gaggaagtgt tgaaggggtga aacttcctgc ttttattggt gaccacagag 60
 tggtagctgn agatatgtct cggggggtcag gagaccttg ggacgtcang tgggggtgcta 120
 ttgccccaaa ccaaacttga ccaatccccga cccaacccgg gcatagtcgg tcagtggagaa 180
 cctgtgatgt acctaagcag gcgagctcct ggcagtcaac agataaaagg aaaacaagac 240
 cacaaagcaa ggaggcttgt ggtggctggc cagctgtgaa tnttgtgtaa tatgtggatt 300
 gtggcctctg gtaatcgatt accanagggtg agtaatcgat tacaaggctt anaaattgag 360
 gacaggaggc taagatggtc tctgngtaat cgataccaag ggggtgtaatc gattac 416

<210> 30112
 <211> 437
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30112

 aaatgtggca tttgtgcaca atacacaatg tcctgtacat tacaacaaaa tgggtgatca 60
 taaaggcgta atagaacttt aatggatatg attaggagta tgttaatcaa ttgacttta 120
 cccgtatctt tgtggatgta tgccttgaaa actgtcatgt atttgttgaa tagggttcct 180
 agtaaggcag ttccaaagac acctttaaac tgtggacaaa taggacacct agtataaggc 240
 acctgcatgt ttgggggttg caggcagaaa taaggattta taatccgcaa gacataaaat 300
 tggatgcaag aacaatcagt ggatatttca ttggttatcc agaaaatgaa aggggtatat 360
 gttttattgt cctaatacata gtatgagact tgtcgaaact aanattgcaa gattcattga 420
 .aaatgaataa atcagag 437

<210> 30113
 <211> 407
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30113

agtttgttca aatcaaatca ctctacatt tgatctctag catgcatctt ctttgtttac 60
ccactcctca cgtntgggtt tttcggggaa aaacaccata actaaacgcg ccgcaaggga 120
tccctatcgc accagatcca aatctagaac gatgggtgat caagaggaga cgcaagaaca 180
gatgacagcc gacatgtcgg ctctgaaaga acaaatggcc atcatgatgg aggccatgtt 240
aagtatgaag cagctcatag agaaaaacgc ggccaccgcc gccgctgcc gttcgggtgc 300
cgaagcagac ccgactctct tggcaactac gcaccatcct cctcaaaca tagtaggacg 360
gggataggac aactggagc acgatggcag cctcacctg tgataca 407

<210> 30114
<211> 301
<212> DNA
<213> Glycine max

<400> 30114
agagcttcat gtttgttctc cttacacctc cattaattgg ctgctttacc ttctcttcca 60
ttgcccgcct cttcattttt ctccatgtat ctctcacat gtcttgtgat aaatgttttt 120
aacatgattc tttagagtct ccaccaatta aacttgctat agaagctaga tttgattttc 180
tatggttcaa atttcttggt cttgttcttg aaccatgaat tgtgttgagc ttaagatcct 240
ttgagttctg ccttgttatt ttctgtggct gagacctaca ccatacaatt cttaccaag 300
t 301

<210> 30115
<211> 352
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30115

atcttactct ctatgagtca cacccttgaa gtaggatcac aatacataag aagaggatga 60
gtcacaccct tgaagtagga tcacaatata taagaagagg atgagtcaca cccttgtttt 120
ttggctttaa aacattgttt tggaacaat ttctaaact gaatagattt tgaatgaaaa 180
attaattcct gaatagtgtg aaattacttt agaaaatagt ttttaaacc aaaaaggtag 240
aaggaaatta aatagatcct aaatattttc tttatgtaca ttntatgatt attatgttca 300
atgtcttcat catttactta gcttggaaata atacaaactt ccactttctg cg 352

<210> 30116
 <211> 370
 <212> DNA
 <213> Glycine max

<400> 30116

ttttcataga aattgcggttc tacctaaggg tgcaaatgaa aattttcttga cactcattcc 60
 tgatgtagag aattcgattg gatcgaatga gtttagaccc atttctcttg tggggtgtct 120
 atacaaaatt gtagctaaaa tactttctat ttgccttacg aaagtgttgc acaagggtcat 180
 tcatgagtga caattggctt tccttgaagg tagaaatatg ttagatggag tggttatagc 240
 aaatgtgtcg aacatggatt tcctaaatta aagggtgcat tgagaccttt actatgtcgg 300
 ttcttgtgaa tgggagtgca actcacgagt ccaaaaacga gacgggctcg gtggtttgta 360
 cgaatgccat 370

<210> 30117
 <211> 388
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30117

agcttttatgt tgctcattga ctccaaaatg ctacaaagaa ggacagagat ctgtatgggtg 60
 atttgcagaa gaacataaac cacagactct tgcaacaggt gcagatttct gattcatggc 120
 aatctgagtt actaagggtga ccaaggcatc aagttttccc tcaggctttn tattttcaat 180
 agatgaagat gaattcgtgg ccacctcatg gactcctcta aggacaatag catcatttct 240
 tgactgaat tgttgggagt tggaagccat cttcttaatc aaattcctag cctcaacagg 300
 ggtcatatca ccatgagctc caccattggc aacatcaacc atactcctct ctatgttgct 360
 aagtccctta tagaaatatt gaagaatg 388

<210> 30118
 <211> 445
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30118

gcttagcgcg tgaagaaatg gtgcttagcg caaggttttc gcttagcgga taagcaatct 60
gaaaatgttt ctaattcatg ttctacttat ctcttcacac ataattttta caaccctttt 120
tggttcattac taaacaagct gaaatcaatc acaatcacaa gcaagatgtc ttaactacat 180
gcaaaaaata aaaatgaaga tagagaaggg aaagaaaagt tgggttgctt cccagtaagc 240
gcttctttta tgctactagc ttgatgcac atcctgttat ccaggatcca ataatgttcc 300
cacttcaagg accttcttct caggtcttct ttctccatc acatgaactt taaaatagac 360
attccggtca agtggctctt tatcttcatg aaatagatca aagctgattc tctgatcttc 420
tatgccaatn tgcaacatct tcctc 445

<210> 30119
<211> 393
<212> DNA
<213> Glycine max

<400> 30119
agcttgtagt tggtgaactt gcgaacctta caagtogaac tcttatcatg gccacccat 60
atgaatgata taggggtgca agaacttgcg ttcaattttt tttttggctg atttatgaat 120
caaatttcat aatataaata aacatttttg agtttagtta caaaacatat tagtttaaac 180
acatttgaaa atagattttc gaaagtgttg aatctacact ttggaaactt agtttctaga 240
agtacaagca ttgttcaa atacaaattag agtaccttac tgaatctcca tgctccatta 300
tgtatgtatt cccctcgta ctaaacctct ttggacccat tgggtctcaca tcaagacacc 360
attgcattga agactcatgg accacccaca tgc 393

<210> 30120
<211> 357
<212> DNA
<213> Glycine max

<400> 30120
aattcatcga aggaaggtgc gaggcaataa ctcaaaaggg ctcaccatct tggatagcag 60
atatggcagt gctcagggcc acgatgttat ttacaatttc tgaagctgct caagctagag 120
agaatcatta catgcacatg cgcatgcacc ttcttacta attacgtacg ggtatagata 180
ttcccatagc taatcataat ttcttggtct ctctttaatt gtgtgtatat ctatttattt 240

ctattacgat gcatatatca tgccatagaa tgatgatctt accgattttc tacaatacat 300
 ttgtcgatca actccgacgg aagaaatcta agaaatgaaa gaaaatatga atatgac 357

<210> 30121
 <211> 414
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30121

agttttattga aacaagctaa aaagagctaa ttttaaaata agttnttcta aatgcccctt 60
 gagtctatctt tgtaatgagt aagtcattaa ttttgtcctc aaactattac cttttctttt 120
 gaatgatggc taatgtaaaa aataaaatat aattactcga gtagtgccac aagtattgta 180
 tatcatgcta agtaatcctc caaacattaa aaaaattggt caaattgatc cctaaattnt 240
 tctcanatac ataaacttaa ggaccaaatt gataaatatt caatactacg gngattagct 300
 aaatactttt attacttttg ggacaattta tgagatagaa agggccatac ttcaatgaca 360
 aaattgatgg tttattcatt gtggaatctt canataaatg gtacagaaaag atat 414

<210> 30122
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30122

tttctctcca ttttttttat gtttaaaatc atctatgttt ctcttanatt gngtaacttt 60
 gtgtttgaaa aaagataaaa aaggagggtt acatgataaa tccaatccac ccccatccc 120
 tacctctttt tcccttctcc atctanacac actagaaact tccagggcac attccaaact 180
 canngcaaaa aaggtgaagc cataggatca catattactt ttgatatacc ccaactaaga 240
 aaaaaatntc acataaacac aatntaaaat tatttttttt caatttcacc tccattagac 300
 tcagtatcgt cttccttccc aataacatat cttggagctc cccaagttcc attcactctc 360
 tgccctcagta tatangactg aattctcggt tgattgtgtg caatctagct tcagcttctc 420
 aggttcttct ta 432

<210> 30123
 <211> 324
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30123

tttcttttat cttctgtgnt ctgggaacct ctcttctctc atgtgtaccc aaaccaatc 60
 acctggctca agaacgactt tctttctgct tttgttggtc tgccttgcat agctcgatt 120
 tttcttttca attagagcct tcacttgctc atgcaacttc ttcacatact cagctctagc 180
 ctgtgcatcc ttatgcttaa acatancaat gttaggcata ggcaacaaat caagaggagt 240
 caaaggatta aatccataca ctatctcaaa tgggtgaacaa ttagatgtgc tatggacagc 300
 ccgattatna gcgacactca catg 324

<210> 30124
 <211> 396
 <212> DNA
 <213> Glycine max

<400> 30124

tctacttatg tggcagggcg ggcttccttc accttcttgt ctccaacgcg aactttgacc 60
 attggtgttc cttcccgcga tgcttctttt catgtccgcc tgagtgggct tatagcctaa 120
 accatacttc ccacgatttc cttgggtatt tatcaggcta gttatgccgc cgttgttttt 180
 tcctaaacct atcccggtt cataaccgtt ccccaacata actcgggcca tcattaccgc 240
 tgcacgggac agacaaggct gcccaaagag ggagtccacg gaggaatgc tgaccacctc 300
 aaaagactgg aaagcagttt ctaacgattc ttctgcggct tccacataag gcatggagga 360
 tgggcagctt accaagatat acttctcgcc tgacac 396

<210> 30125
 <211> 345
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30125

accaaacc aa acaacattca acccaaccct aagttcaact tcattctttt ttcatattgat 60
 tcgctcccaa acagagacat agcctctttc ttttcgtttg agaaacaaac ccctcgtttc 120

aatcattgat ccttttctgt taggtttgtg aatntgcttt tgtttttgta aaactttgca 180
 cctccccctt tttggatttc gtagttaggc gaaaatttta atgtttccgt gtttcaaatt 240
 tgcagatacc agttactctt ccaatttcgt catggccaaa acctccttca agcttcagca 300
 tcctttgggt acttnttttt cccctttaat atttcgttnt ctcta 345

<210> 30126
 <211> 443
 <212> DNA
 <213> Glycine max

<400> 30126

gaatgaaata cagtgtacag actctaccct actgtttttg caagttaaaa caattgcact 60
 agagaaatat tgccaaaaga agcttcctag ctagatctcc taaaaagcag ggatattata 120
 gaagttggaa ataccacccc tacaaccttt ctttgatttg atcctcttac tgtgtccttc 180
 attcctatgt tatggagcag atccccctcc aaagaaaaaa gaaagaaaag cattcaaaca 240
 agttttgata gaaaagtcgt agagctagta catacaagta aagctaagaa tgtaggactt 300
 tgattccttt gctttttgtc tctctttctc tctccccatt aaacggaaat taaattaaaa 360
 aaagatTTTT tttttcattt ataaaaacaa aaggctgata aagctgaatt catatacaca 420
 gagcagtttt acagtcggac atg 443

<210> 30127
 <211> 273
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30127

acaaaagatg atcttctacc cccattccta gttatatgct anttttcatc gtatattatc 60
 ggaggtataa atctttaatt ccaccttgat tttgatatta cattaaccat aactcgatgc 120
 tagtatatac gaaaaaaaaa ctgtaatttg attacttacc ttatgctcta atagccagag 180
 gatcaaggct tctctcacct acgcttgctt attgtcttct agcatatgca acaaatgtaa 240
 ggacagacac ttttggaatt catgtacttc atc 273

<210> 30128

<211> 419
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30128

agcttctggt cattgggccc gtaaaactca ctagggatag cgggtattatt aaacgcagac 60
 aaagcaacca gcgataaaac ccaatacaga taaagcacct aaactaatag aaaagtaagc 120
 ttctccagac cataccagtg cagcccgagc ccatgcaaaa ggtttggtta agatatgcca 180
 gattccacca agtatacaaa tggaacccaa ccatacatgc cccccaatta tatcttccaa 240
 atcgcacaca ctaacaatcc acccttttcc cccaaaaggt gattttaata aatatccaaa 300
 tataatactc ggactaatgg tcacattggt tatttttctt acatctccgc ccccgaggagc 360
 ccacgtatca tatatnacct ccaaatanag agccttgaat actagatgaa acgcaccta 419

<210> 30129
 <211> 495
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30129

agggaccgcg agtgagnctg agaccttgca caacatagtt atttagcatc catccccgag 60
 agggaaacag cccggatcac cagctaattgt ttttaattga cgcctcagtg acaaaggagg 120
 taggggtgca gagacaacca ggagggttgc ctataagcag acacccttga aagagtgcgt 180
 aataactcac tgatcgagcg ctcttgcgcc gaagatgaac ggggctaagc gatctgccga 240
 agctgtggga tgtaaaaaag aatccgtagg ggagcgtctc cgcttagagg gaacgaccg 300
 cgcgagcatt gctggacgac acggaagcga caatgtcggc ttgagtaacg cacacgttgg 360
 tgagaatcct atgcctcgaa aaccaaggg ctctctcgta aggttctctc accgagggtg 420
 agtcatggcc taagatcatg ccgaaaggcg tatcgatgga cacaggcgaa tattctgtac 480
 tacctttgtc ggtcg 495

<210> 30130
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30130

tttatatgan gatanagctc tcttgcttct gaattcctta ccaaaatcct ttgaacattt 60
caaggatgcc attctttatg gccaaaaaca aaaacttacc cttaaagaaa tcccgacctt 120
caccaggac cagggaaatc caaaaccgcc aggggttctaa tctgaggata atgggtgaaag 180
cctgaatatt ttcaaggaaa ggagtgaaaa aagggaacaa gaggaaaaag tccatatcaa 240
gtcaagggat tcaaagaatg gctagaanac aaagttcana tgctttaatt tgtcacaac 300
tggtcatttc aagaaagact gcccatacaa gatcaagaaa ggatctttgg actctgctga 360
catagttgaa gcctctgang gtatgagagt cangtgttta gtagcttcta tcn 413

<210> 30131
<211> 445
<212> DNA
<213> Glycine max

<400> 30131

aagagcccag gtagtcgaag agaagttcaa gtccatatcc ttcaaagtct gaaaagagta 60
tgatgaacta agagacgtca atatggccac cgctgaagcc ttggaacgag aaaccaagaa 120
ggcccgaag gaagaacacg accaaagcaa agttttgagg ggctttatag ggcagcaata 180
gtgagctcaa gctccgaaga ggtgaaagga atcatcatgg gtcaaaggca tgatcttgaa 240
ggacgagcta aaagcttgcc tcatgtcgaa aagaaatttg tcccaacagt taagcgagac 300
agaagggaat atgtgggcca tcatcgatga gtgcaaagag aagctaaatc tatcggcgac 360
tcataagcaa aggctagagg atgagtacgc caagatatca gcagacaggg aagcaaggga 420
tagggttatt gattcattgc accaa 445

<210> 30132
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30132

agcttgatg attatggggt acccatcata tgtggtacta ggtggcaatc aggcgatggt 60
gcaagtcgac tctccacatc caciaatcac acataaatcc accatcccca gttgtccacc 120

ttcaactgag ctcacgtgct cccacgtagc ccttatactc gttcctctca acaccgggct 180
 cccatcaatc cctccaagct tccacaacat ccaagaaatt cagcatccaa acatcatgaa 240
 ctatccaaaa ccaagaaaac agggcatagg cagaaaactc ttcccaaaac acattccaat 300
 accacagttt tcttactca nataccccag taacattctc tttgtttcga ttcgttaacc 360
 ggtggatcaa ctcanaattn ttactggagg tccctaatac atatatctac ag 412

<210> 30133
 <211> 411
 <212> DNA
 <213> Glycine max
 <400> 30133

gaataatggc ctcacaaat tatttatttc ccgaagttat tttctataaa taagtctcct 60
 attgttagtg gtgtgggtta ccattattgg aacaaccaca tgcaaatttt tatagagaga 120
 attcttcttc ttcttcttct tcttcttatt catgagattg attaacggat cgagggtttc 180
 ttaagttgaa ggaattctga acacaaggga agggttgtgc ctatgtgggt cagactttgt 240
 aaaaggcatt ttacaagata gtgaacatct caaacgggtt gtttggagat tagacgtacg 300
 cacagggcat gaccgaacta gtataataac tgagtttgca ttctctcttc ccttaaaatt 360
 ctcttactta ttgggtcttta tcttttgcac tacagaagtt tactttgaat t 411

<210> 30134
 <211> 353
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30134

gctttgtttt aaccanatat gaacaattta tgcgaatcat tctttcttgg aaaactttct 60
 ataaattctt gtaagattaa agctctcaaa acacctttta taccttgaga aaaaagactt 120
 aaagtgttga gtgttatatt tgtctataag accatcacta aaattaatcc atgtgtaatc 180
 ttttaacaaa tctttgtgat ttgtttaaag ccaacaatgg cttgatagaa caaagaatat 240
 tggtttaaat cacacttggc gtgagcttgt acgtgaagag ctagaagtga cagtgaataa 300
 tacttgaac tctgataagc tagtggaac ttggttgta ccaagaattg aat 353

<210> 30135
 <211> 300
 <212> DNA
 <213> Glycine max

<400> 30135

gcacaattac cccttccact cttccatata aaatatctac ttttaaccagc tttgcttccc 60
 tgggtgcgtc taaataaagg attcaagaaa tgttgtgcct tcaacaggcc aacctagaag 120
 aaaccatggc atagagcttc ctcttctaata gcatggaaac caattcgctt atcaagaaca 180
 ccttggaagc ttcaactcaag tacttcaacg actctcgtat agttctatat taatatgcta 240
 atgcaacaac gacttgcttc aaccggctaa tcccttgtgc tacaaccagc tacttctttc 300

<210> 30136
 <211> 357
 <212> DNA
 <213> Glycine max

<400> 30136

agctacttga tgttgaatcc agaatagatt agagagtctt gatgatcaca aagatgatga 60
 caaagagccc atgagaatga gttcaagatt gacatctgaa cacttcaaga atcaagagga 120
 aatttgagtt caagattcac gaatcacggt tcaggattta agtttcaagg aatcagagaa 180
 tcagcgagtc aagaataatc gagttgaaga ttcaagagtc acgtgaagac tcgattcaga 240
 taagtacaca aacgtttttc aaaacattga gtagcacatg aatatctgac aaaacctgtt 300
 gccaaagagt ttttactctc tggtaagtga ttaccagatt attagaagtc gatacca 357

<210> 30137
 <211> 383
 <212> DNA
 <213> Glycine max

<400> 30137

agttttctct cttaaatttc tataaatagg gggagaagtg aagtataaaa gggttcagcc 60
 ccttaagcac ttcaactctc ctcgaaatag ctgacgaaaa ttagtttcgt gaagaaaatc 120
 caagccgagg cgcttccgta acgtttccgt gagtaattac gcgaagatgc tcgaccgttc 180
 ttcaagattc atcattcggt ctctcgtttc ttcaagtctc aacgggtaag tacctcaaac 240
 caagcttttc aattcattct atgtaccggt ggtggcccac atttcgcttc atgtattttt 300

attctcgttt tcattacttc ttatacccoct tttagcgtgc ttaagccatt tattttaagtc 360
 atttctcgct ttaatctaaa aat 383

<210> 30138
 <211> 500
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30138

cgancacggc cntattgaca ctnttagatt ccccgccact cgtagatta ttcaacctcg 60
 ccaccagct ngactatgca ggctacgggtg ctacttctat attctccgcc ttctggcgga 120
 acctgctgga atgctcaagt gggcctggct cctatcctca ccacatgat actaaatata 180
 cccaaaccac ttactcgttg attcctcatc cgtaaccgta cggaactcta tgaatctcgc 240
 aacgatactc gctctatgac cagaatgtca cgaaacctta cggattacac aatcatacct 300
 tatttggcct ccgaactgta actgaacttt accgactgag caacaatgct ctcttttgac 360
 gtaatgcac gcacccaact ctacggatta tacaacaccg catcctttcg acttctgcga 420
 tgtcacgaaa ctttaccgac ttactcataa tgggogcaca gcaccttcaa gcggtcaacc 480
 atggtctctc ccccaaacc 500

<210> 30139
 <211> 411
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30139

ttgcttctta caagagacta agaaatttct gacaaaaaat cttgagatga aagatcttgg 60
 ggaagcctct tctgaattag gaatcaagat actaagagat cgctcttaag gtatcctaag 120
 gttttcacia gagagttata tcgataaggt cctaaataga ttcgacatga aagatagtaa 180
 accaggagat accctgatag cttaaaggaga caaatttatt ctcaaacaat gtcccaataa 240
 tgaccttgaa agaatagaga tgcaaaagat tccttatgca tcaacagtag gaagtcta 300
 gtacgctcaa gtttgcactc gtnccgatag agcatttgta gtaggaagtc tgggcagata 360
 tttgagtaat ccttgaatgc agcattttaa agcagcnaaa cgtgtgatgc g 411

<210> 30140
 <211> 409
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30140

ggctctagcc tcaactcaccg cctttctggt tttattttcta gctatcttat acttatccgg 60
 agtttgagaa tttctacacc tagaccactc cttgaaacac tcctttttta ctctaacttt 120
 gctctgaaca ttttcattcc accaccacga ttctttaccc ctaggtccaa aacctctaga 180
 ttcacccaac gtctcttttag ccactttaat aatctcttgg gacatcttgt tccacatata 240
 atttgcaatt ccttgtgatt gtccacacca accctcccat atctnttggt ggaagattcc 300
 ttgtttctca cccttcaagt gccaccattt gatccttggg gctaccatag gacttcttct 360
 ctttgcccta tctctaattc ttacactcat aaccaacact ctatgttgg 409

<210> 30141
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30141

ttgagaagat tgggggttgac ttacctatgg aactattaca gttggcggtt atgagttatg 60
 ctgaatgtca tanagtagtt ggagacctgg accaaaataa gatagctttg gaaatttttag 120
 ctgttcctga ccttcctcaa ttggctccat tttttctaag gaaatcatca cccacggca 180
 atgaagacat tgtgggcca ggtattcctt ttctgttct acttgtgctt aatgaaattc 240
 acaacgggta ctcaaatttg gaaggagacg cactttcagt ataagcagag cttggcctca 300
 aataccaaga agttatgc 318

<210> 30142
 <211> 351
 <212> DNA
 <213> Glycine max

<400> 30142

gtgaccagga tcagggtcca cgatgggaaa gctgtatata cagaaattgc gcaggaataa 60

gcgaagctcg cccaacttg aagtaaaaag cacttcacag aaaactaaag aaactccagc 120
 atacaaagcc ctagaagaga tactaaaaac catgcctatg aacagaatat gcagtatata 180
 atagaagcaa taaagaaaac ttgtcacta ttcacaccaa taacaacaaa catagtccca 240
 gactcaaaat acacctacct tcaaacaaat agaaatagaa atatagacac agtacaatgt 300
 tatcattggc accattctaa catatagaga ataccgcact gaatctacac a 351

<210> 30143
 <211> 447
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30143

cgcccatgtg tcatactatc tgggcaatca gctcgtcccc gngatccttt aaatcaacct 60
 gcaggcttgc aaccttgact tgtctgccta agcacactat gcctctggaa gttttctttg 120
 aattaagatt aacctaaactt ctggctcttag cccttggtgg gtggtgaggg aggttaacct 180
 aacccccctt cacccttaac ttaacttttt tattggattt taaagttttg cagttaagct 240
 aaatgcccc tgtgcgctaa cctggatgta ttctgataac gtgactaagc gcccatgcta 300
 cactaagctc actctcttta ttgaaaaatg ggacctggct aactcacttg ctgcctaact 360
 taattacaan aaatatttgt gattcagcta tgcagttact ggcttatcct agaaaattta 420
 aagcgcgcta cgcactgctc ctaacac 447

<210> 30144
 <211> 432
 <212> DNA
 <213> Glycine max

<400> 30144

ataaaagtta ttatgttaga catttgtgag acaaaatgat caagaggcaa tgggaccctg 60
 aaagtgtgaa gttgagagaa acctagatga agtgtaggct actttatgag tggcgtagt 120
 ctatgctcaa gtccttggaa agtggttatt gtgtgtggaa ctgtatgggt catgttggat 180
 caagtcgaga atctagaagg ggggtgaata gattatttca aaatcttgtg ttgtcaccac 240
 aatctgttgc ccttgcactt tagcacacaa gaaccagta tcaccatcaa tatgagttat 300

ggtatagaaa aattttaciaa ggtctatgta gtaggtgcac ttcattctcca ccaattttctt 360
 caaccctgta taatactagt tcaactcagg aaaaaaaaaa tcttcacttt tgagccacgc 420
 caaactaaca tc 432

<210> 30145
 <211> 344
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30145

ttaatgggtat atttaccttt attagccata ttaaataatta tttattatta aacttaaatt 60
 taagccaatg gtaatactaa aaattgggtat tttttaattt ccttaaattt ggaaaattcc 120
 ccccccccc ttggagaaat ttcctaattc tgtccttgca atcagaccaa gtccagtgtc 180
 tggagtagat gaacaagtgc tcaatcttgt actgtatgaa gaaaaggatg aagacacatc 240
 agataccttt agagtatgtt tggatgggga aacttaaaat tctgagaaat ttaaattcta 300
 gaatttcnat acttcaatga attctttatt tcaaaatttt tggt 344

<210> 30146
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30146

tgagactttt attactatat tttccacttc ttatctaacc ttggaaaggc tccacaaaga 60
 gaaagccaat attagaaata tgtttatctc tgatgaatgg atcctaaaca agttatctaa 120
 ggagcctaag gggaaagaag ttgcaaaggt agtgctcatg ccttcttttt ggaatagtgt 180
 gggtttacact cttaaagtca tggctccact tgtcaaagtg attcttcttg tggatgggtga 240
 aaggaaacca gccatgggct atatttatga agcaatggac aaggaaaaag aaacaattat 300
 caagtctttc aacgacaatg aaagcaagta caaagatgtg tttgcaatca ttgataaana 360
 gatggaattg tcagcttcat aggccattgc atgcatctac ccacttctta 410

<210> 30147
 <211> 211
 <212> DNA

<213> Glycine max

<400> 30147

tttttcgaac catttccgtg aataataatt ttttggccaa atgggccaac aggcaatttt 60
cgcccaataa atgggaaaaa gccatgttcg gcccgaaacaaaagcgggt gggctcgcac 120
aaaagaaact aacccgacta cattttaaat tttgtatgca acacaaaaac aagaaaactt 180
cctgtgccgt aaaaaaaaaa cattacatga c 211

<210> 30148

<211> 435

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30148

atgacgccga tcgaacattt cctaaccgac gtcttgctna tttcggtcag ggattgaatt 60
gaaaactcgt taggcgacat ctgtcgcgaa gtaccgaccg atatttttca gccgacattg 120
cacaattctt tttagaaaag ctgcgtggtc gataatggtc tttttacggc agagtaagtt 180
ttcttgtttt ggtgttgcatt aaaaaagtta caatgtactt cggctagggt tttcgtgcga 240
gttcaaccga cattttgttt cgccaggaa aacattagcc cacctctgca aaaaaaatat 300
ttgctaaccg tcttcatgca tatttcattc aacgattgaa tagaaaactc aatagccgac 360
aacggtcgtg aaatagtccc gactgatatt tttcagccgg cattgcgcatt ttttttctaa 420
aaaaacgctc gctgg 435

<210> 30149

<211> 168

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30149

atgcttcaat ggaggaaaat aaagagggag agaaagagag aggggggggg ggggaacacg 60
aaattgaagg gaataaaaag ggagagaagg gaactttgaa gtatgtctca caagactctc 120
attcatcaaa gttacaacaa gtgttacaca tgctttctatn tatagact 168

<210> 30150

<211> 445
 <212> DNA
 <213> Glycine max

<400> 30150

acgggcatct tagttcattc cttatgaata atgatttttt tagaggaaaa tggatacaac 60
 tatgttatgc aagaattatg attcccaatt tagcaatttt attaagaatt ggcttccacg 120
 ttttctctct tcttgggtta gctccaatag ggataccaag atacacaaag ggaaatttat 180
 tgatcatata gtttatgata ctagcatacc tctccaaagt gctatcttta accctaatag 240
 tcctaaagaa acttttatga aaattaactt taagtcccaa gatgagctcg aaacctctta 300
 atatactttt aatgggtatac acatttgaga gggatgcac accaaaaaat aataaagtat 360
 catcgggtata ttgaaggaga ttgatttgag ttttttcctt acgcaccaag aagctacaaa 420
 acaaattttt ctcaatagct tgtct 445

<210> 30151
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30151

agctnttgat caattcanat ggtcataact tttaactcag atgtctgatt catgcgcata 60
 atatatcgag acgctcgaaa ttgaacaatg gaagctcttg agcaattcaa atggtcataa 120
 cttttaactc agatgtctga ttcaggcgca taatatatcg agacgctcta nattgaacaa 180
 cggaagctct caagtaattc aaatgggtcat aacttttcac tcggaggtcc gattcangcg 240
 cataatatat caagtcgctc gaaattgaac aacggaagct ctcgagaaat tccaatggcc 300
 atcttttcac tcgnggtcc gatttaggcg cataatatat cgagacgctc ganaatgaat 360
 agcggaagct ctcgagaaat tcacatggtc ataactcttc actcgagagt ccaattc 417

<210> 30152
 <211> 446
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30152

gggcaacaat ggtggaggaa gataagaaga agaatgtttc tgtgagagag agggagagct 60
 tcagaatttc ttttggtctga gtgaggagag agaacagctt ttggttttta aaagggtttt 120
 ctctttttct attattctat tcaagctatg ccacatgtct ccatttgagt ggagcgaana 180
 gggcccactt tctcttttga ttgtgactca tactcagcca caaaaagtga gaaaatctga 240
 cctttgaaac gctaaaatcc tgectcggtt tgcgtgtcat ttctctgggt ccagttcctc 300
 gtgtttctct gcgtccgtcg gngccagttt tcgaaagtag gcaatatata tatatcanaa 360
 cgctcanaat aaaaccccga gcgtgggttca gaggttggtt ttgttaaatt ctaagtcgca 420
 cgcaaatga tgatctttta actaat 446

<210> 30153
 <211> 392
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30153

agtttttgtt tcctagaaaa ccaatatattt tttagcccag cctcattaca agccaataaa 60
 agtccttctg attcaatttg tgtatttctg actttatggc atgagatgat gtacaaagat 120
 tggacctctt gttagtgtt attggtaaatt aacttanaca cttatgcgtg agtgatatag 180
 tggccgtgag aatttggtta aatatctttc catgaaatct gtctcttgcc tagcttcatt 240
 tagttgtgtt gttgactaac atgttctttt ctctgaanaa ctgcatgtct tgtgaaaagc 300
 aattgataaa angcattttg gttcatttgg tatcatgtaa ttaaaatttt gtgaatcaca 360
 cacctttgta cataatcact gcatgtttt ca 392

<210> 30154
 <211> 444
 <212> DNA
 <213> Glycine max
 <400> 30154

tcttccacct ctctgcaatt agaaatccac tatttctaaa ggcgttgaaa acctggtgac 60
 acattaaact tatcacaact aaataccgtg agagagtcca accctttcag caatacttcc 120
 tctcccaaaa attgagcaca ttcagtaatt tcaagattgg aaaggcatgg gaacatgttt 180
 tccccatact cccttgataa ccttttaaag tttggatgat gacggatggt cagatcttct 240

agagccctga aaactatttc cccatcacag gactcctcat agaggatttc tacgtaatta 300
 ttattttattg tacccaaaat ctttaaagaa ggcagtttcc ccaatagtgg aagttgtaaa 360
 cagttttcac aattcagcag atttaacagg gttaaatact tgagagaagg agtagacatc 420
 cattgtggga aatgagcacc ttg 444

<210> 30155
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30155

agcttgatg tattatgggg taccatcac atgtggtact aggtggagg cggcgatgg 60
 tgcacaacaa attttccaca tccacaaatc gtgcataaac ccaccatccc ctgttgccca 120
 cctccaattg agctcacgta ctcccacgta gcccatatcc tcgtttctct caacaccggg 180
 tccccatcaa tcttcccaag cttccccaac atccaagtaa ttcaacattc aaacaacaca 240
 aactatcaca gccaaagaaa tagggcaaag gcagaaaact cttgccaaaa caccaaccaa 300
 aatcacagct tttctcactt aaagacccca gtaacagttc cttcgttcca gttcgttaac 360
 cgttggatcg actcgaaaat ntactggaa gtctctagta cataatccta cat 413

<210> 30156
 <211> 441
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30156

tgggacatct tgacttgctt tccaatctga cattctcctt atattctgcc ttcttctatt 60
 gtcagattgg gaatgcctct aacagcacct ttgtcaatga ttttcttcat gcctcttaag 120
 tgcagatgtc caaatctttg atgccatatt ttgacttcat cttcttttga gaatagacat 180
 gtggaggagt aactggtttc ttgagggtgc cataggtaac agttgtcctt tgatctgctg 240
 cctttcatta ggacttcaact cttctcattt gtcaccaagc attctgactt tgtgaagttt 300
 acattgaatc cttcatcaca caactgactg atgctgatca agttcgagc agtcccttc 360
 accagcagta ctttgttcag actaggaagt ccatcatgga ctagctntcc cattccagtg 420

atcttttcctt tagagccatc t

441

<210> 30157
<211> 410
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30157

agcttttttc aattctgcca ttttaagggt caaattgatc ttggaagtca aaacctttgc 60
acttgaataa ttggactgcg agtttgggct tttgttcgtg taattaattt aactagttta 120
attgggctgc gaagtttgtg caacttggtg tccaaagttt atccccctatt ctgagtgaaa 180
gtaacctctt tgggggttaag tttgagttaa aattgccaaa ttctgcctct atgagtttta 240
tcggtatggg caatttggtc atttcaaagg aaattatctc agaatgggct aaaactttgc 300
caaaatgtag aanaattcat acatcgaggt gcccctgtga gggacaaaca caaacattan 360
gaatcattnt tgccaaattc atttatctgg accacttttg gaattccttt 410

<210> 30158
<211> 424
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30158

ttcttttgtg ttaggtatta agatactacg agattgctct tttgtatcct aaggttgtca 60
caagagagct atatcaataa agtacttgat agattcgaca tgaaagatag taaaccaggc 120
gataccccaa tagctaaagg agacaaattt agtctcaaac aatgccccaa taatgacctt 180
gaaagaactg agatgcagaa gattccctat gcgtcgtagt agcaagtctg atgtatgctt 240
aagtttgtac tcatccccgac atagcatttg tcgtaggagt tctgggcaaa tacttgagta 300
atcctggatt gcagcattgg aaggcagtga aacgcgtaat gcgttacttg aagagaacaa 360
aaggctacat gctcacttat tagaagtntg acaatntgga gatcatcggg tactcagact 420
ctga 424

<210> 30159
<211> 348

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30159

 agcttggtgt atgggttcaa ttcctaacta tgtatagaga aaaaaagtca tttgttgaca 60
 aaggttgat tcaactttatt ggtgaaagag tgtagtctgt agctggtggt ctgaaatac 120
 ttttaataata cacaaagtat gtatttaca aaaaaaatg atacaatttt cattgctaaa 180
 gacgggtgta agacaatcat gaaaataggg tgcaggggaa aaaactaaag ctcagatcga 240
 gcagaataga gcagggcagc agcttcttat ttgattgat ttgtcgggtt tcatttattt 300
 taaaatgtaa tttggagact ctatgtttct ttccttntc ttttatga 348

<210> 30160
 <211> 96
 <212> DNA
 <213> Glycine max

 <400> 30160

 tcttttgatc cgactatgtg actaatcatt gattcttgtc ttattcaaac ttaaagctca 60
 tctctcgttt gtaatagtgt atcatgttgg gatgac 96

<210> 30161
 <211> 391
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30161

 agttttgagc aatttctaac aacaataact ttttactcgg atgttgatgt gatccttacg 60
 gngcggatcg cttgatacag gctgtagagt tttggatgac gccacttcca gtgaaggaag 120
 ataagtcagg gtagacacca cttccggtga aggaagataa gtctgggcag acgccacaag 180
 gattaccttg ataagtctga gattggttca accaggaacc cagagagaaa ctcaccatat 240
 tctatcatat gccagaagct ttgtcttatt cagaacgaaa accaataactt atagtgtagc 300
 tgaacaacaa gataaaaata gacatgggcc ttctaaacag tttgggccaa aattacaata 360
 aaaataaatt ataactanaa acttatttaa c 391

<210> 30162
 <211> 439
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30162

acacagcaac acagaatcta ggtgtccaac actccttcaa ttcaatgggt tttctaggtt 60
 tgagaagtga aatttagaat gaggtaaatt tggagcaaac tctcacctca cacaagtcta 120
 taacatcaat ctaaacttgc tcaaactgga tttacaccta aaattccacc gaatcaaaat 180
 ttgactcttc aacacccaaa tttgccctag aaatggctct ttgttcactt tggtcatttg 240
 tttttccctc tatcacagcc taacctttct cataagtcct aaatggcatt tcaagctaag 300
 attaactcgc tctaacctct aaatactacc aaatccagat ttggccttcc agccctcaaa 360
 aattcactct ntttccactc ataacaccac attntcactt tctaacccta ggtaattct 420
 accattcatc tctaacagt 439

<210> 30163
 <211> 404
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30163

ggcgggtctcg atgatacgcg gagatacctt acggnatacc gcaccctttt gtcattccaga 60
 ggcgggcgggc ccgatgacaa gcagagacca agtttgggtca ttctgcaccc ttgtatcatc 120
 caaaggcgggc gggcccgatg atacgcggag ataccttacg gttatccgca cccttttgtc 180
 atccagaggc ggcgagcccg atgacaagca gagaccanac ttgggtcattc tgcacccttg 240
 tatcatccag agggcgggcggg ccgatgata cgcggaataa cccgagtggg tattcgtata 300
 aacattcttt tgctatctgt aagacagaac gctngatagc atgcagaggc tgacatagtc 360
 ttctgcacct tttggctcct cggaacaac aagtcattta catg 404

<210> 30164
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 30164

catgatgaat cttgactatc tacacaataa ggtgctacat tccatgctct tttcaagntt 60
ttgctaccta aagccgcatg ccaattcaag tatattttcc ttgctgact aaaattgtat 120
tcaaattaaa ggtatacat ttttttgtaa tgtattttct ttacataaca tgcaacatat 180
ttatgtatat ttttttgtag gacattttga ctaccaaaaa ttatatgcac atacatccaa 240
gtattttgct atcataccca aagtgtaaat tgccaaaggt attttgctac ctattctaaa 300
cctacacatt catgacgagc aaaattccta aacatctang cgtanggaaa ttattgtagc 360
gtggcccata gctgattgct ggccaaaaag ggtaactnta cccaatatng cacctctttt 420
gtgtctttt 429

<210> 30165

<211> 412

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30165

tagcttttta tgaaggaaat atgcttcccg actcaacgta tgactatcaa aagattatta 60
aggatcttgg acttgattat gttaagacag atgtttgcat tgatgattgt atcttatata 120
aaggaagcta taaaaacctt gatgaatata ctatttgtaa gaaaactaga ttgcaagaaa 180
ataagaagaa aaataatgtc cccaataaca cagttcggtg ctttccaata aaaccaagac 240
tgcaaaaatt gtttaggtct aaacaagtta tgtaataatt ttggtctcaa cattttggaa 300
aatgaaaggc tccttcagtt ttggaattga gaagacanaa tgaatggcta ctaatgagtg 360
gtaatgacca ctaatgggtg gtaatgacca ctaatgagtg gaatgactac ta 412

<210> 30166

<211> 440

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30166

aagatttgaa acttcacata taagagatgc tntattatta ttangttaaa ttaatgatga 60
tctcaaaata aaaggtgaga ttgttttagca acttatgaac ttgaaaattt tgagtttttg 120

ttaagtacga ctatttggtg tgatcatatta tttcatgtaa actccattag taaaaagtta 180
 caatcaaaag atatgggtat gtatgttgct atagaacaat tgaaaggtct tatttctttt 240
 tttgaaaaat atagagaaga ttgatttgaa aatgctataa tttctgctaa agaaattggt 300
 attgaaatag atatagaacc taagcttttt gaaaaatgtg ttattcatag aaagaaaaac 360
 aatttgatga gaatattgat aataaagttg taaaattgcc taaagaatca tttanaattg 420
 attacttctt gtatataata 440

<210> 30167
 <211> 415
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30167

gcttggtgcta tttctagttc atataccata cctttaagcc aaaatgcttc cttcactcct 60
 tcaactaggg cccttgattc tgctttaata aatgaaagaa caaccactga atgggtgaatt 120
 gctttccaac taattggtat acccaacaaa agaaatacat atccggtaaa gactttcttg 180
 tacctacatt ttctacaaaa tctgcatcta catagcctgt gattgctgcc tcatgtgttg 240
 tcttcttgta ccttaatcca gcattcaaag atccatttag ataccttagt gtccacttca 300
 cagcttccca atgtgcactg ccaagatctc ccatgaatct gcttataata cttacagcat 360
 gagccaagtc aggtctgctg canaccattc catacattat gcttgcaaca ccact 415

<210> 30168
 <211> 435
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30168

tggaatcgat aaggcgggaat gtctgaggta tcttattacc ttccatggtg ttgaggataa 60
 tttgggaaat gagtgaaggg tcacggtcac ggggtgatgtc ggggccagca atgtcttcat 120
 ccgcaagaag caggtttaaa tgggacttgc aacgggtggcc gtgagaccat ttatcatcgc 180
 aatggtaaca caggccctgg tctcagcgaa ttgcaagctc ttctgccgat aaacatttga 240
 cgggattttt ggtagagggg agggaaagtg tattataagg tttaggggaa aagggtgagg 300

gggtcgtggg ataagggaaa gggtttgggg aggatgtggg ggcacgggtt cctctgcagt 360
 ggtccaacat tntatcccc tggagacgtg ccagctcaat tgcctgtgga agcgagatgt 420
 ggcgcaacac atgga 435

<210> 30169
 <211> 413
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30169

agcttgttgt atgcatggct agagatattt gtgtcgactg aagaccaaga atttgtgtata 60
 tatataagtt ggaagattct gtggaaaccc aattgccaaa gaaagtgact gtccatggat 120
 ggagggcata cgcattgagat atcacatggt gttgtcttcg gtccaactat gaaggataag 180
 ttgtgaagac ttgcgtaacc ncaccttatt atacattaat tggccagcat aaagtctcca 240
 catgtaacca aatgagacgt gcctctgtga aacgtaacgt ctattcaagc attatttggg 300
 attccatatt tgatattgtt tttgcgatcc aagtcacac accttcactg ttgggaattg 360
 caaagaattg tctgtggagg gagaaatatg catcgacga agcattacaa aat 413

<210> 30170
 <211> 441
 <212> DNA
 <213> Glycine max
 <400> 30170

tgtttctctc agggacaccc ctttcttttc caatctcatt gttcctccac aagctcccct 60
 ctctccctat ccaataccag agagaaagat gaaccacacc acccacttcc aatcacacca 120
 ttgtttcctt caaaaccatc acttttcttc ttggtgacat cttcatcaat gccaccaccc 180
 cataaaaaat cttccatttc catgcctcaa taacaataat aaagttaatg ttcaaaaggg 240
 taggtgattt cacaaatagc acgtgaaatg ggaaatgggg ttgttttgtt ctttcttata 300
 gtgggttttg agtgtgtgtg tgtttggaac taagtaagaa tgggtgagtgt ctgaaacagt 360
 gtaacagtgt cagtgcagc atatgaagat ggcaaaagaa gtaataatga gaaaagatct 420
 gaaggggtact aagtaagaat t 441

<210> 30171
 <211> 414
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30171

agcttggttat tatatacaaa acacacaaat tattatgaac aaattgacgt taatgacgta 60
 aattattagt aacacttacc actgcatgtc tcaactaagt cacatcagac ctttcagaag 120
 tcgaccgtgc tgctggctcc gtgaaccgac ggatatctgt gtctggatcc tgaggggcaa 180
 ctctgggctg cgtagcatga ccatctgccc gaggatctga tggctggccc ggtgtcatga 240
 atggatgcga aatgcggaag aatcagtcga tgtagtcgct tgcacactgc cttgcacaa 300
 cgcagatgtc acctgctaca accatatggt ccgaatagtg catccacctg ttgtgtatat 360
 catcagactg caccatgaa tcngcangtg gagcangaat ggtctgagtg tate 414

<210> 30172
 <211> 400
 <212> DNA
 <213> Glycine max

<400> 30172

taccogtcac atgtgggagc aggttggtgg actggcgatg gtgcaagtcg actagtgaca 60
 tgcacaaatc acacatgaat ccaccatccc cagttgccc ctttcaactg agctcacgta 120
 ctcccacgct agccttatcc gagttactct caacaccggg tccccatcaa tccctccaag 180
 cttccgtaac atccaagcaa tttcaacatc caaacatcct gaactatcaa aaggaagcac 240
 atacagggca gacgcacatt actctgccc acacacaggc caataccact actattatta 300
 ctgtataaac ctattaacac taccttatgc acaattgggt caccgggtga tcaactcgaa 360
 gatttactgg aggtccctag cacataagtc tacatttga 400

<210> 30173
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30173

agcttctctg cacaaatatt ccgtccccta aatatatgcg taaatttaaa attaaaagta 60

tgaattttat atgtatatag tttaatcatc tttcctaag ttttcaaagc accaatatta 120
gagttgattt aaaatatattc agtactaaaa gattatttac acttatattt aaaatgtcat 180
tttaacgtaa tgtgttttaa ttttttaaatt attatagttg taataaaaaat aattacacat 240
ttattntaag tatgttattt taatgtaata tttcttaact ttttgtggt tcatnttttg 300
gtttataatg tggttaacta ttatttgtga attttgttct tgagtcttat gtctaataaa 360
tataaaccaa ttttcatgtt ttcttcaaaa cattgtatct gtctcttttc ttt 413

<210> 30174
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30174

acatatcaaa tatgttcatt nttccatctt cttagacttc tgctaatagat ttcctttcct 60
atgttctgtg cttttttggt gcttttaggtg gtccttcatt ccctactttt aactctgtca 120
aattgagatg tcaatgccat aaataagtta acagtaacag caatgtgcag ttcttcaccc 180
cataatcagt cacagtgcac ttccttgtct catgggatat tattaaaagt cagtatctta 240
aggtttgagt actttgggta ttgcaatga atctgoccaa ggtagagtag caatgggtctt 300
taaggggctaa ggagatataa taatcaaggg gaaatataca aaacaatcaa gaggaactag 360
taaaaatata acaatatata caatgaacca agtaagaatc tc 402

<210> 30175
<211> 409
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30175

agcttctaca ttcaatttca agcttttcga tatattacgg gactcaatcg gacatccgag 60
taaaaagtta ttggagtttg aatttgctca gggcttcggt attccatttc gagcgtctcg 120
atatattacg ggactcaatc ggacatcaga gtaaaaagtt attggtgttt gaaattgctc 180
agagcttcgg tattccattt cgagcatctc gatatattac gggactcaat cagacatccg 240
agtaaaaagt tattgtagtt tcaatttgcg canggcttcg gtattccatt tcgagcgtct 300

cgatgtatta cgggactcaa tcagacatcc gagtaanaag ttattgtcgt ttgaatttgc 360
tcagagcttc tacatttcac ntcgagcttn tcgatatatt .acgggactc 409

<210> 30176
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30176

attcanacga caataacttt ttactcggat gtttgattga gtcccgtaat atatcgagac 60
gctcgaaatt gaataccgaa gcgctgagca aattcaaaca acaataactt ttactcggga 120
tgtctgattg agtcccgtaa tatatcgaaa agctcgaatg tgaatgtaga agctctgagc 180
aaattcaaac gacaataact ttttactcgg atgtctgatt gagtcccgta atatatcgag 240
atgctcgaaa tggaataccg aagctctgag caaattcaaa caataataac ttttactcgg 300
gatgtccgat tgagtcccg aatatatcgg aacgctcgaa attgaatgct gaagctctga 360
gcaaattcaa acgacaataa cactttactc ggatgtctga ttgagtcccg taatatatcg 420
agacgctcga aattg 435

<210> 30177
<211> 342
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30177

agtttgttgg tcttctcata aagatttata ccttggttgg aattcttgac aggttgactt 60
ctttgctcct tggtatactg gttccccgaa tgattcctcc agcttgagcc ctgtgtgaaa 120
ttctccctt gattgaatct cagcggtcct cctaggttgt agccttgga tctgtggcgg 180
ttctgagctc ccatgtagtt cacctccatg taagaatcta cttgngctat tgattgcctt 240
gtttcatgtg ctccatcaca gatatggcat cccctatnt gcatgagtga agagtgagag 300
ggacttaccg cttgtaaag ttgagggagc ttgctgaggg tc 342

<210> 30178
<211> 447

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30178

 cagcaaatgt cttcacaaat aatcatcaca cagcttattt ctagcaagac taccatcat 60
 atctcccaaa accccatacc cacgaaaatc aagagggaaa gaagtccacc caaacctgaa 120
 atttcgaagt cccactcgta gccacacact tcacgactcc aaaaacgccc tcttttcacg 180
 atttggggca gaaatgatgg ccaaagggtg aagctttgct tggagcttca atggagaatg 240
 aagaagaaga aaatggcaac gtgagggaga gagagagctg tctgaaaagt gtgggggctg 300
 agtgaagaga gagaaaagct ttttggtttt aaataaaagg gtttttctct ttttctatta 360
 ttttattcaa gctctgccac atgtccctat ttgagtggag cagaaggacc cactttcnct 420
 ttntactgtg acccactc agccaca 447

<210> 30179
 <211> 379
 <212> DNA
 <213> Glycine max

 <400> 30179

 gtactcgaag ggggtgaccc acaactatat gtacgattca gaagtatcct catcatttta 60
 ttaagtgaag cgatgatgtt cagaggcgga ctcatgaggg taccacttgt ctccatcagc 120
 atccatacat tgattacaag tgtgagtata ccccgctttc acctgcctta acgatcgta 180
 tgagcctcat ctgcctgact aatatgtgat gacaagagac gaccctacta tgggtcgatc 240
 tcggatgcta gcaccaatth gtatgcacga ccgtattagt agtcaacgtg tcgtatcgcc 300
 tacttgcaac gtagccaccc ccaaatacat atgtgcggag gagatgtcct ccgccgctga 360
 cccaccacag aatgctgcc 379

<210> 30180
 <211> 338
 <212> DNA
 <213> Glycine max

 <400> 30180

 tttgactgcc agaagctaca aacaattcta ttgcatagac acacacgtgc tatgtagcta 60

caatttcaat ctctggagac gcttaaaccct tcagcagtat atagcaatta gatatacaaa 120
 cactatcact agactttgca tacatcgaag caccctatat aaagtaaaca aacttgctaa 180
 agcaaggaac atggatcgat ctgaattaat tcccacaata aatacccaca agagaaacag 240
 aggctcgac gtcacatcc acggccatgt gacctctctc tctctgctgc atggatatat 300
 ctttaccatg gagccggaca tcaagcatat attgcca 338

<210> 30181
 <211> 351
 <212> DNA
 <213> Glycine max

<400> 30181
 acctcgcccc gggatctcta agtcagctgc aagctgtttt tttttcttgc taaacggcta 60
 ttgtgataaa atgttgata ataatgctct cattttctgt gtgtaattaa ggtcctccgt 120
 ccattcctcc ttcgaacgtt gacctcatat gttgaaatag gtttgccacc aaaaaaggaa 180
 accattctca acagtagcat gtcgcatatg cagaatcaat ttataaggc attactgcat 240
 aaggatctgt acgtcgcgaa tgcacgagga gaacgtaact tcttctaaat atagcaatgc 300
 tactacctaa atgctgccat caccocatatc tcttacacgg tgctgaacct g 351

<210> 30182
 <211> 252
 <212> DNA
 <213> Glycine max

<400> 30182
 tgaccttaaa gccactattg agtgatttat gttgatacag cgtcattttt aacatttatt 60
 tcttaagata ctgctgatag atctatcgaa ataagatggt tttagccttt cttttctata 120
 ttatcttaac cgtcaataat ttactctatt tattgtatca aactattttg ttgctgtta 180
 ctcagattct atcctataaa tgttattcta gttgaaattt gtacgtatta tctcaattac 240
 tcgatttttt ta 252

<210> 30183
 <211> 302
 <212> DNA
 <213> Glycine max

<400> 30183

gcaggatgct agcttgtagg attatggagc acccatcaca tgtggtacta tgtggtggtc 60

cggcgatagt gcacaacaag gctttcacat tcacaaatcg cgcctaaact cccattccc 120

tgttggtccac cttcaactga actcacgtac tcccacatag tccatattgct cgtttctttc 180

aacaccgggt ccccattaat tctcccaagc ttccccaaca tccaggtaat acaacattca 240

aacagcacia actatcacia ccaagaagac aaggcaaagc cataaaactc ttgccaaaac 300

ac 302

<210> 30184

<211> 379

<212> DNA

<213> Glycine max

<400> 30184

cgtgggaacc agaggtgggt aatataatga agtgaccaag atcaatcaga aatcataacc 60

aaccaaaaac ataaataagt gataaccaaa atgaaatcca aacagtcact attcagaacc 120

acatagaata aaaacatata agactaaagt ccaaatacta aaagataaat aatgtgctga 180

aagcaataat caaaatatca tagccaaaat acacgactta taagacacat agaattataa 240

actaaattct aacaagggtg aggtgggtgg ggaagatcga aactctgacg aatgtaaccc 300

acatcttctt caagctgtgt gaggcgaata tccattccgg caaagcgagt atccagtga 360

tcgaaacggt caccaacat 379

<210> 30185

<211> 388

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30185

tagcttgtgt ttgtctctca tggctgatgt caaatctatg acatgtggat taaactcaac 60

tttaattgca aaaaaaata attaaaggaa tatatattag tgaaacttca tcatttagaa 120

tataacttag tttttctaac atttttctct tctgctccat taaatctctt ttacatttgt 180

ctttttctct tttatattgc ctttatcttg taccctccca cttcttctct aagttaaaat 240

taagacaaga atagaaacta gaaagggtag agtttggatt tttgcaccat atatgcatga 300

tgcccttatt gacatgaaat cttcattntc accatatgtc taggtcacca aacaaatctt 360
aatattggtg ttttttttta ctgggaca 388

<210> 30186
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30186

ttatcttggg ttatctggag gtgtaagaca ctggtttttt ttcattgctta tcttanatta 60
tgtataaatg gtcaatttca ataataaag tataaccttg ttcattcagg attgtctaga 120
aaacatcgat gcagttcctt catacaattt tgtgtcatat attttgatgt tatttgttat 180
acaaagatcc ctcttaagct taatggaaaa tttaactgtg caatcatatg atcaatggtg 240
ctttaattat ggtgctgaat gttgggcaat taagggctaa caagagcata agatgtggtg 300
cgcagatgaa atgttgcatt ggatgagggg tcacactaga aaagataaga gaaagaattt 360
attaagagag aaattgtgag ttgcttttat acaggatatc aggcctataa tagcttggac 420
acaagacaag aataccaata 440

<210> 30187
<211> 349
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30187

cacaacaaa ttttcacatt caccaattgc gcataaacct accaatccct ggtggccacc 60
ttcaactgag cttaagtact tccacgtaac ccataatctt cgttctttta acaccgggtt 120
cccaataatc cttccaaagc ttcccaacat ccaagtaatt caacatttca acaacacaaa 180
ctatcacagc caagaaaaca gggcaaaggc aaaaaactct tgccaaaaca ccaacaaaaa 240
tcacagcttt tctcacttaa agaccccagt aacaattcct tcatttcagg ttcgtaaccg 300
gttgatngac tcanatattt cactggaagt ctctagtaca taaacctac 349

<210> 30188
<211> 395

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30188

 tcgagatgag gaagtgtaaa agggtaaaac ttcttgcntt tattctgttg accatagagt 60
 ggtacctgga gatatgtcgc gngggtcagg agaccttgag gacgtcaagt ggggtgggtat 120
 tgcccaaaac caagcttgac caatcccgac ccaacccggg catagtcagt cagtgagaac 180
 ctgtgatgta cctaagcagg caagctcctg gcagtcaaca gataaaagga acaaagacca 240
 caaagcaagg aggcttgtgt ggtggctggc cagctgtgaa tcttgtgtga tataatgggtt 300
 atggcctctg gtaatcgatt actaagggtg ggtaatcgat tacaatgctt ataatgaag 360
 acaggaggct aagatgggtc cttggtatcg attac 395

<210> 30189
 <211> 413
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30189

 cagctcgtcc cgggatctct aagtcaccga ggctgcagct tttattataa tggttttgac 60
 ataaattaca tatctggaaa atctctatga cttttgaacc tgccgctggc cctgtcctac 120
 acttaaatta ttcttctctt aatgcctggc attttttctg gtagaaaagt gggttttgac 180
 cttacttcct tttgctatct aactggggct tagttgaaaa aggggcacat tacacattct 240
 taaagttaag tgatttagnt ttctacatct gtatgtgact atgtgtggac taagggtgtt 300
 gatgtactaa tgtacttctg ctgtcatcct catcctggca cataccttgt gttggtacat 360
 gattatatta ctagcatctt agatgcctat aggcatgtga ttcccatcaa tta 413

<210> 30190
 <211> 451
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30190

 tcatgctatt gtaaggtnnt tagcacanat tccaaatgag tggtgtgata atttcatata 60

gaactatata tcagaatgtc attaaaaaaaa aaaaccaata caaacttcct caagactccc 120
 cttaatatgt cattcataag acttttgaat gtagcagagg cattagtgag cctaaatggc 180
 ataactagtc attcataatg tccatggtga gtcctgaaag ctattttata cctatcctca 240
 ggtttcaaca aaatctggtg ataaccagac cttaagtcca acttggagaa aaattcagct 300
 ccaaacagct catcaatcaa ttcatcaact gttggaatag gaaatgtatc tttaaccgta 360
 atagaattca atgctctata gtcagtgcac accctccaag aaccatccct cttcttaacc 420
 aanataattg gagaagaana tgggctctta c 451

<210> 30191
 <211> 391
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30191

actaacctag aattaaaata acttaatggc attaacctan ggaattaaaa caaaattaat 60
 ggctgagtgt aactgaaatt ggggcaacca aaagtcaccc ccaacagccc acaagtcagc 120
 caccatttgg tcttccaaaa ggctgatgcc tangttgcc attgggccct tattacaact 180
 tgaactacat cccttttagt tgattaaccc aaaacatatt tttggtcagc caactttaca 240
 aggattgggc cattatttag acaaactaaa cactctaaaa ttgaaacaaa gtgggtgcat 300
 ttagtcctcc tccatttggg ccatgatata actcacaacc ttggactttt ctccttgaaa 360
 cttgggcttg tattcaaata gtatggacaa c 391

<210> 30192
 <211> 431
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30192

cgacaataac ttttgactcg gatgttctat ttgtcccgta gtatctcgag acgctagaaa 60
 ttcaaaacag aagctattag aaaaatcaaa cgacgataac tttttacacg gatgtcccat 120
 tgagtcccat aatatatcga gacgctcgaa attgaaaaca atagcactta gcaaattcaa 180
 acgacaataa gttttgactc ggatgtccga ttgtgtcccg tagtatatcg agacgctcga 240

attgaaaaca gaaactgtga gcaatttcaa acgacaataa ctttatactc ggatgtccga 300
 ttgagtcgcg taatatatcg agtcgctcgt aaatgaaaaa agaagctttg aggaaattaa 360
 gacgacaata acttttgact cggatgtccg attgtgtccc gtagtatctc gagacgtca 420
 naattcaaaa c 431

<210> 30193
 <211> 395
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30193

actgaaatta aacactgaaa cataaatatt aacccaatt ataaaatgta cttaaaacca 60
 aataataatc aaagtggta aaagaacaga aaataaaaat tctatcatgg gtcctgtggt 120
 gcaaaagggg catcatgtgg tgcagaaagg gcataatcca tggcttgggc atcatcctca 180
 tcctcagata gctctagcac aggcgtagcc accgtcgatg cttgcaaaga agacaactcc 240
 agcacagggt tggctactgg taatgcttgt ggagtcattc ctagcgaatc cttcacagt 300
 tccttctgag cagttggatc aatctcttgg atgtctggct ctttaataact angtaacct 360
 ctacaacatc tggatcatcc ttctgagtag cttct 395

<210> 30194
 <211> 447
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30194

tctgatagtt tcatgctntg atgatttgag cttctgcaa atggtcttgt tgttaatttc 60
 gcttgatggg tgagatattt gatgcttgct ttggaccctt ggaagaagat tgagaaaacc 120
 gatgttggtga gaagatttct agaagctatg taccatgggt gcacctacat tntgatttgt 180
 ttctatattt ttgttttggc cgatgtattt ttgacatggt tagctatatt ccaatttttc 240
 agattttatt ctcatttcct ttaatgttga tgtatatctt ctccaacttc cagcttgctg 300
 gggaggggga tattagagat atagattagt ttagttagtt acaagttagt tactagtatc 360
 aattatataa ggtacaatgt atttatgtaa tgagagagtt ttgctcattt gagcattact 420

ccaatattaa ttagttctac cttttcc

447

<210> 30195
<211> 404
<212> DNA
<213> Glycine max

<400> 30195

tctgtttatc ataattgttc ttcagaatgg agaacttcag aacaaactca aacctatcat 60
gtcaagtaca actagcacac tttcatgttt tgcaagaagt tgtcacagag ttaaaggcct 120
tacttcaagc aactattttc ttgggatatt ttagacttgg aggaagctat taaagttaga 180
aaaaaaagaa gacctagttg tcataatggc tactttggtc tgtgggtcct atgaattgat 240
agagcatagg tgcaaaactt gagagagaag gtaggagata acctacactg tgtagaaacc 300
atattcatct tctttccgta ctacatatgt gctatgtagc ttagaatggg tgtggcaaga 360
gtgcatatag ctgagagctt taaagaagat tcaccctaata tagt 404

<210> 30196
<211> 404
<212> DNA
<213> Glycine max

<400> 30196

ttcgtcaccg cagcaacact gtagaaacct tactgcttcc tttatcgaca cctactgggt 60
cttttccgac taccgacttc ccctttatct cagttagtaa ttgtttaatt tcaactgaat 120
gtggctttta attgaaacaa ttaacacaga ggtagcagag aagaagaaga caaatttggg 180
tttatttgca cgcctagta gtagcagaca cagagaagaa gataaatttg cgtttgtctt 240
tgaaagggtta aatattgtat gttatataga agtactaaca ttgtgaaaat ggggtcccata 300
tttttcaagt ttatctaata tctcatatgg tgaaattata atattgtatc atactgatca 360
cttcaataaa tgtcattata aaaaatatgt actttataat atat 404

<210> 30197
<211> 395
<212> DNA
<213> Glycine max

<400> 30197

agcttctata gttagtcaca tcataatacc atttgctttg ttgaacgtta ttgtaggaac 60
 caaatcaacc aatggtgttc aagcaaaaga aaatgttagt tgagcagatt atatgcagca 120
 ggcacatcgc ctcatTTTTac cgcctactga agacaataaa agtggttaca ggaggaatga 180
 aagatggaag aagccattac caaagccaaa ttaaaagcaa actttggcat tggaaccatt 240
 attaggacca ttaggggtgaa tttttgacag cagcaacaaa gtttatgcca aatcttatag 300
 tatcagagtg gaagaatctc tggccttaag atggtgtatt gaagtcatgt aagaaaggac 360
 caaactgca ttacacatat tgtgatggca ttatc 395

<210> 30198
 <211> 431
 <212> DNA
 <213> Glycine max

<400> 30198
 ttagccatga tatcttgatt ggaacataat tttttcttct tttgcttgtg tgctctaagc 60
 acaagaacta aatcttaata atgaagaagt atgttaatct taacttattt accttggaga 120
 atttgaattt tcttgaaata caacaaatta tactagaaga ggggttgaat aatgtgtag 180
 tcaaaatata aaatattttg gaagtgaagag atgttataat agacaagttt atagaaccat 240
 tgtctagtga caaaagggtga actatcatcc aatgagatgt aaaactttgt ctttcagtaa 300
 aaatcttgtg tgtaagggtg gacaataatg gacaatgaaa tattttataa gttaatgaaa 360
 aacagtaaaa atactttgtt ttgtactagt tcaactcaacc taagttatgt ccaattttcc 420
 tttaccaact a 431

<210> 30199
 <211> 381
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30199

agcttgtatg atgtcgagcg tactgatgcg taccatgaga tgtctgcggg ggtttaaccc 60
 acatgggctc tgccataccc taatttcgtc cggggacctt tgcttgatga catgcgacct 120
 ttctttggct cttgtgaggt gcttggcatc catcattagg caatntgtga aattccagga 180
 cataccgaan aaccaaaaaa atattgatgc acaatccgta agtttccgtg acaccccgga 240

aatcaaattgg aagcatcgtt gcataattaa gtgaggttcc gtaacattcc gtaagtcaaa 300
 aggggggatga ttatgtaatc cgcaaggttc cgtaacatta cggaaagaaa acaagtatcg 360
 ttacgaaatt cgtaagtttc c 381

<210> 30200
 <211> 439
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30200

tattcaattg cttcagattg ttgcacagaa tgtgtttatg tctgtgtggt ggtcgacaga 60
 ggagcataaa ccacaaagtc tggcgacagg tgcaaatttt tgattcacgg ccagttgggt 120
 taccaggta accaaggcat ctagtttacc ttcaagcttc ttagtctcgg ctaatggaga 180
 tgaattcgtg gctacttcat gcactcctct aatgacaata acatcacttc tagtactaaa 240
 ttgttgggag ttggaagcca tcttctgatg gaagcttgct tgtgggggctt ctatggaggc 300
 tggatctttg agcttcaatg gggctccttta atgggtgattt tccaccatgg agatgcagtg 360
 gaagacaaag gagaagaggt gagaggaggc gccatccact anggaataag ccatggaaga 420
 aggagcttca ccaccaaga 439

<210> 30201
 <211> 421
 <212> DNA
 <213> Glycine max
 <400> 30201

agcttctttg agcaatgaca aacaaattcc acacaaacaa cataaccttc tagagacaca 60
 ttaaagagag tcatcactag atctctctac ctctatcttt agttaaagca aattgttcct 120
 atgttagata tatttgagaca acatagaggc agctatcttt ccttacctag aaaaaaagaa 180
 aaaagatgaa gtttaaacta cagataaacc acatgtctaa aattagtttt tagcataatt 240
 taaaatagaa aactatatat tattccgatt gtaagcaaaa tgagtcttta gagcctcaag 300
 gcactacaac acaggcacia taaattttaac ataaaaattc acaataaaaa ttggcttcaa 360
 agccaagaaa gaacaaagga aaaaaaagaa caaagtttca gcaatcatatc ttgggagtat 420

<210> 30202
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30202

gagattataa taggattgta tggttnttag gtattgttta ctctcaatac catatgcagt 60
 ttctcaagga ggggtggtga gtttcattct gctaattggt cttgcaatga tgttttggta 120
 cacgnggtta cttctacaga ggtgtatgaa caagcatcca ctaatcaaatt cttaccctga 180
 cataggtgag gttgcatttg ggctaagagg aagagctatg atctctacat tcatatacat 240
 agaattgttt ttagtggccg ttgagcttct gatattggaa ggcgacaatc tagannaatt 300
 gtttcctcat atgaacttca naattggtag ccttagaatt gaaggtaaaa gtggttttgt 360
 ggtgctagct gctntggcca tactaccaac aacattggtg agaagttngg agctttggct 420
 atgtttctc 429

<210> 30203
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30203

agcttttttag ctttaagaac tttttccttt ntacatgccc aactctttga gtgacatttg 60
 tattgattat tgcattcttag tctttatctt ttcatatgta catcatgcat catcatgtag 120
 aggtaagaag attgtttcta aagttaaaaa aattntcaat gcataaaact ctctgttnta 180
 atcaattaca aggctaatacg taatcaatta cacaagtgtt ttagcttgc agagatattc 240
 tagtttcagt ttaatcgatt actagttaac cataattgat tacataaatt agttgagatc 300
 atgtttgatt tttcacgagt ctctgtttta atcgattact agatgatcat aatcgattac 360
 tacattctta aaggtgttcc cagaagtgat ngagaactct ttaatcgatt acatcaa 417

<210> 30204
 <211> 391
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30204

ttatcntttt atattggtat accatgctac agccgctccg gccaaagctgt cttgaaagaa 60
atggaccaat aacttttctg ccgcagaata cgccctcatc tttcggcaat acattcgaag 120
atgccctttc ggacatgtcg tccctttgta cttatcaaag tctggtactt tgaacttagg 180
agggatgacg atgttgggca cgagacataa gtttgctaga tccgagaatg ggtaatttcc 240
gaggcccttt accgctctca gcctctcctt aagcgcatac atctttccct tctcctctgc 300
gaagggaaca tattcgatta cgggtgcggg tgaagatggg acgtggcgga ctatgtntgg 360
ttgngtagt tcatgnggg atggatcttt g 391

<210> 30205

<211> 363

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30205

tgaaaggata atattccggg taagtttttg ctattaaatg tcgcgtacct ggaactattg 60
gaaggatgtc ttatatatgg cataagagca attttggtc aagtaaagggt gtgaagttga 120
gtctagggttt gtaaattaag aaaattaaat aatatactcg accaaatgca ggaaatatta 180
cccgtattgt agaaaaaag ttccttaaaa aaagtcggtt cactatagta cgtacatata 240
antttgatat atttaattta gtgaatttct ctggacttga actcataact agccagatga 300
agtagataat ttttttttan atcaaattca caataaaaca agaaaaatct ctccatgact 360
tgc 363

<210> 30206

<211> 407

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30206

tagcttgtat gttagagtag taccacagga cggtgataaa acaaaatact aaagacacat 60
aattgggaca ttttactcaa ctatctttga taactcttcc tatcccgagg atatgataat 120

ggagcatctc aataacaaat aataatggca tatgatagac ataagtgcac gcatgctaata 180
aatttaataa tttcatgttt taatgtgcac attgataatt gtgtgattag taaaatcaga 240
tacagttgta aactttcaca ctctgactca tgagcaccct cattcccact atttaattga 300
tagatcccct ctaacaaact gtctataact atntgtcact tcccttctat cttanatgag 360
atggctcatc gtctcccccc ccccccttca tgctctagag gatacaa 407

<210> 30207
<211> 417
<212> DNA
<213> Glycine max

<400> 30207

gcttgaagaa gtttgacttt actatcctaa ctccccttgag tggcattcgt attggttggt 60
atcttgtatg ttgcatctta gtacatatga tatcgtattg catcatgtat catcatgggt 120
agtgtaaaga aaagtttctt caagaggcaa aaaatctttg ttttaatcga ttataggttc 180
attgtaattg attacgacat gttgtctaaa gcttgaagag ttgagtctca tatcggttta 240
atcgattaca ttggtgtttg agacaatgat tgatttattc aagagtctct actttaatcg 300
attacgaagc ggattaatcg attacttctc gctcgtctag tagttcaaaa gtgaacaaaa 360
acactttaat cgattactta gagcatctaa tcgattacat tgttcttgag ttatttt 417

<210> 30208
<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30208

gagagatgaa ggaccacgat ccagacagtt cgagagattt gcggagcgaa gatttgcaga 60
gaccagagcg cgaacaggaa gccgccctga gagccagaga tgaggctgag agcgactgag 120
aggccctaga cgcggaagag acatccccac aactagtacg acggcaaacc gtcaacctct 180
acactcccgg ttgcaaagga agcagactag ctatggaaag ccaaatactc tgctggatct 240
cccttgacga tactngatgg aaatagcagc atatctagac aacgacaagc gcatgatcac 300
tgagctatca gaacagcata ccgccatgct actgcctaga ccacgtagat gcatgggctg 360

cgaagattat acaccagtgc aaaccg

386

<210> 30209
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30209

cttctacaga atgatgaaat ggattcggct cttttatnat gttcttatgc agntttgatc 60
tgcagaatgt tttgctattn tgggtgttgat acccacgagg aagttgttgg ttttgttgag 120
gtagagtttg aagtgaagac caaagttttg aggcagatgg tgtacatcaa gtctgaagaa 180
gctgggttta atggattgag aatagtagaa aagatcaaac agaacaagaa gatacagaaa 240
ataagccctt aagcctcttt gagcagatga tgtagcaaa gatgcatgag ctcatgagga 300
tacatgaaga agactatgct aagcttaagg agtgctctga gtatattgta gaagcaagct 360
tcatgatgaa tcaagattga ttcanggagt tttgatgatg acaaagatga tngacataag 420
ctcacaagta aagatcactt catgat 446

<210> 30210
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30210

tgaaatcgaa ataaggatat ttgcgactca atctaaaaaa taagggagag ntgtctgtta 60
cgcgatgact ggcccgggac ctaaacaacc gaggcacaac caggtcttta gctccacaat 120
gcaccagggg gaatcatatg caacaatatc cgacgccatg gaatgaccaa gacaaactct 180
aaccaccaa acaccttttc taaatgtaca aagaccaaac tacccaaccc tgcctctcat 240
agctacgcta ccaggataac taatacacta gaaaccaaag acaacgaaaa caagcccaag 300
tacgaaaaag ggaccttgca tagaacgaac aatgcaccaa taaacccatc cacgaacgaa 360
atgatataat gaaaacacaa gaaacccacc attcccaata cgaagcaagg aaacaagcac 420
ataatctaac gagaaatacc aaaagcaacg 450

<210> 30211

<211> 490
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30211

 gaacctaact atttaagaat ttcattataa anaannnnnn nnngaggaat gatctctaac 60
 acaanaannn gnnnggaggn aagnaacgaa gcagagcacc ttgtttatct tctgcaacta 120
 cctgaaggat tttttgtgat gacagtatac aatgttaccg acttgacttt gagtgccatg 180
 gaatgataaa gatccaactt tgtattcacg gtggaataaa tgggtgtaat gagaaaaaat 240
 tgattgggtt gaacaactgt agttttattgt tgagatactg tcaatggtct atggatattc 300
 ttgatgcttg gctccattgg ggcctcaggg gaagataata gatgaagctg ggaaagtgtc 360
 cactatggag ctaaataata tgcctagttt tgaacacgcg atcaatctgc tataggttat 420
 ttgttgcaac tgagaaagag aaatgtggtg acaagggtta ccttagatca ctactactgc 480
 atagtattgg 490

<210> 30212
 <211> 566
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30212

 atacaacgga gtagtaagaa acagtaaaca cagtgaaaaa gcacgagaga tanantntna 60
 annannntnn nnnnnnnnnn cnaannnnnn nngaaggaag gactgagtcg tagtanntnc 120
 gncannnnncn nnaannnnnaa nnnannnnnaa nggnnnagaa nnaaaaagaa naaaaaaaag 180
 agaaaganga aantatttat agattatgag taaagaaaag aagaagagga aggaggagga 240
 gagtatagaa tagaaaaaat gaaaaaatga gaagaganga taaaagaaaa ggaaatgaag 300
 gaatagaaag aaaaaaagag aaggagagaa gggaataaag aaagaaagaa agaagggagg 360
 agagaaaaga atgaaaaagg aaaaaaagga aaaaaaaga aaaaaaaaag aaaaaaaaaa 420
 aaaggaagaa agagagaaaa aaaaagagag aaaagtaaag agaaaagaaa agaaaaagaa 480
 aaaaaagaaa agaggaaaaa aagaagagg aaaagagaag agaaaaaaa aaaagaaaag 540
 ataaaaagaa agaagaaaaa aaaaaa 566

<210> 30213
 <211> 408
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30213

agcttatgac aatttgaaat tctcgagagc ttctgaagat taattntgag cgtctcgata 60
 tattataagt cttaatcgga cctacgtgtg ataagttatg aaccatttga atttttgaga 120
 gattccggtg gttaatttcg agcgtctcga tatattatgc gcctgaattt gacttgcctg 180
 tgaaagggtta tgaccatttg aatntctcaa gagcttccgt tattcaattt cgagcttctc 240
 tatatgtgat gcgcctaaat tggacatccg ggataaaagt tatgaccatt tgaatttctc 300
 anaagggttcg gtagttcaat ttcgagcatc tcgatatatt attcgctga atctgacatc 360
 cgtgtaaaaa gttatgacta ttttagttta tcgggagctt ccgttttc 408

<210> 30214
 <211> 316
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30214

gtgtatggac catatcgtag ccaattgtgc tcatcgataa tggntccagt ttaaactgta 60
 tgcctaagag cactttggag atattaccat tcaatgcttc ccacctaag ccgatttcaa 120
 tgatggatcg tgccttctac agaaccgcc gagaagttaa gggagatatt gatctccac 180
 tacagatagg cctcacacc tgtcagggtta cttccaaat aatggatatt aacccccctt 240
 acaactgtct gttgaggcgt ccgtggatcc tctcagtggt agttgttcac tctacactcc 300
 accaaatggt gaaatt 316

<210> 30215
 <211> 302
 <212> DNA
 <213> Glycine max

 <400> 30215

ttgtgtagga tggatctagg atatcggtt aaatactcat gcaaataac ctttcgtctt 60

caacactaaa ttagggaaaa ctttctattc atcttgccca attaagaaga aacctccaat 120
aaccgaccaa tgatgatagg aggaaaaaga gtgtggcctt ggcagaaaca tcgaacacaa 180
ctttactcac gatggagtga gtccgacccg tgtcatgatg tgaacaaggc cgatgtgggt 240
gagatatgga tagatgccca tgggggtgtg gcttggattg ggttgggtgg ggcattttga 300
tg 302

<210> 30216
<211> 392
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30216

aaaaccatta aataaaagct gagtgacaaa atattaaaaa tactttaatt tatttaccaa 60
tgcttttctt attgaaatta gtagaaagca ctcccatat gtcagtgact tcaaaaaaat 120
ggaaccacat aaagaaaatg agagtaattt tggatcttta tctacctata ccaattggat 180
tgacattatt caataattta aagttactaa aaaggttcta ttcaagacct ttntccactt 240
caatagactt ccttggtata aatataagaa aaataactga ttacatagt cgacatcggt 300
taattatatt aatcttatta aagagtaatt ttntcaact attctgtatg gaacttntat 360
tatgtatata aaaatcatta aactaatact tc 392

<210> 30217
<211> 325
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30217

agcttttatt ctaattgcta agcgacagct tattcgtggc taagcgtgac ctattatcgc 60
caagcacaat tccttatggc cataattgag gtccatgacg ctaagtgccca gtcatggcag 120
ctaagcgaga ttcattgtgg taatatgagt gctaagcgag tccctctcat ctaagcgcat 180
gctcctctgt acttaagatg catcatttta gctaagctgg ccagagcctg ncttagcgac 240
agttgcaact tttctaactt gtagaccttg ctaagcggaa gaatcaatgc gctaagctaa 300
gccttttctc ccanaaaaaa aactt 325

<210> 30218
 <211> 345
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30218

 agcttgngtt gngctaata gngaaangan ngaccaaagg gaagacaaga gccatatcta 60
 cggtaaattg cgtgttgacg ggtcaaatat tgattcggcg gagttctagt tgtaaaacca 120
 gttcatgaaa gtttacatta atgttataga cttgtgtgag atgagagttt gtcctaaaat 180
 taccctattc tcattttcac ttctcaaacc ttgaaatcca ctagattgac gggttttata 240
 tacctacatt ttgagttgct ttggctcgaa gcttgtctct gggtttacata tgatttatac 300
 atgactaacg acttgtagga tccaatctac gaaaatatgg atgat 345

<210> 30219
 <211> 98
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30219

 tcatgcgtag ctaccatgcg tttaagggca ccaataactg ccttaccata atacgcatcg 60
 ccatgcactc atctgagtag tgatgtactc attaaagcn 98

<210> 30220
 <211> 384
 <212> DNA
 <213> Glycine max

 <400> 30220

 cactggccat cgtttacaac gtcttgactg ggataaccct gtcattacac cgcttaaatcg 60
 ccttgctgca catacccctt tcaccagctg gcgtaataac gaagaagctc gcaccaatca 120
 cccttcccaa catgtgcgca ttctgaatgg ctaatggcgc ctgatgcgag atcttctcct 180
 tgctcatctg tgctggattt cacaccgcat atggagcact ctactacta tctgctctga 240
 tgccgcatat ttgatccaga ccggacactc cgcgacatcc tgtgacgcga atcctgtggg 300
 gaagcaggca tttaaattgc gatattgtga gcgtatataa gattaaatat accgtattct 360

atttgtgaga tatgaaggat aatg

384

<210> 30221
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30221

gtgcngatat gagaggtgag cgtgtctatn tatkataact ctgcactgga cccgatgtgag 60
ctatgcaacg accataggag aaatgagtgc gagaaatgag acgatacatt tactgccgac 120
tatgctatgc gctacactga gtacaagact ggaatgggta tgaccatatt caatgtcgat 180
cggaccgttc tgttgtcact ttcaatcgtc tgtagttatg aggcgctcta atatggtcac 240
actggatata tgttgtgacc atctcaatag atccattatg ccggagcacc atgaacgata 300
cacatgatat aataatactg cgaatctgac gcactctgca taaggtaaga ctcatctaga 360
tatatgagag cgcccatatg actatgtaca gcgattgatt cagagat 407

<210> 30222
<211> 496
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30222

aggcgttcat ttctctgnac acancncann natnatcgtn cccgggatac actagagngg 60
ancngcatgc atgcatgcaa actttattgt gtttcaacac ccagcgncaa aggggggaggt 120
cctataattg catatacttc ctccccccac gaacctagca ttttccgcac aaaccatcta 180
tggaaaaaag atcatattaa actacaatcg ctaacacaac aatgggtgtga attgattcac 240
ataacacggc gattcgcgaa agttgcagag ttctggaaaa cctgtataaa cacgatcatg 300
tgcggttagaa cggcacacac gtgtatcatg aagtttaa atccttgtat acgcacttct 360
gatataggcc catgatgaca agcttatttg gcatcagttc tatatgaact ggtggaagaa 420
atgcgtgctc attgcaaaga agcggcacta ttattgcgtc tgtctattgc cacgtgaaca 480
tattgggttaa gatacg 496

<210> 30223

<211> 419
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30223

cgcattgctat ctttcttggt ccatacctgc acacgcgaac atttggaag ttagtctttg 60
 tgggacatat actcttaagc agaaatggca tataacctcc tcccataaat acaaacatca 120
 atgtatattt agagcaagct tatgtgcatg tttccttacg aacgttcact tgcggaagat. 180
 atcctattaa ccgaaaaaat gcacccatat acaatcaagg cagctntggt agctagatta 240
 ttacacgta cttccaaggt gtatttggtta ctacatcaca cacatctcct tggctaaatt 300
 cacatacatg catactccaa gcatttgggg taccaaaaat tgcacatgtg cacatcttgg 360
 tatttctaata acctatacat acacgaactt catgatgaat cttgactatc tacacaata 419

<210> 30224
 <211> 360
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30224

tttctcgtc tccaccctat gcanacccgc cagtgccgc taatagcgtg tctcgtcaa 60
 cgtaccggca aggaattaca accatatatg agaccaccg aagaaactac ctctccgaat 120
 ccaaacttcc ttcgcttgta gtactgcctt aattcccaa attcgggtct gtgctatttc 180
 ttttttatta ttgtactttc ttctcagatc ccggaggcct cttccctccc tgtctcaaga 240
 gaattccccc gattttctcg agaaagtga acggaattat gagctttcag gacatctaag 300
 ccgggcgccc cttcggttcg aggcgcgggt ngatgaatgg gaagcaagac cccacgcaag 360

<210> 30225
 <211> 430
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30225

agctttgttt aatttggttt gacaataact ntatacagg atgtccggtt gaggccgta 60
 atatctcgag acgcctcaaa tttagatccg aagctctgag aaaaattgaa ttgacaataa 120

ctttatacac ggatgtccag ttgagtcccg taatatatcg agacgctgca nattgaaaac 180
 ggaagctcgt atgaaattca cagacaata actntntact cggatgttcg attgaatcgg 240
 gtaatatatc gagacgctca aaattgagac tagaagctct gagcaaattg atatgacaat 300
 aactctatac acggatgtcc ggttgagtcc cgtaatatat cgagacgctc ccaattgaaa 360
 cggagactct tatgaattca aacgacaata actttttact cggatgcccc acagagtgtc 420
 gtaatttatc 430

<210> 30226
 <211> 495
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30226

ccccaccan ngggaattca gtcangacat cgnnaataac atcgaccccg aactttgatg 60
 acctgaggca tgcaagctta cttctttntt agtaatgacc cactanccta gaggggaatat 120
 acttaatggc cttaacccta ggcattgaaa aaaactttat ggctgagtgt aacttanact 180
 tggtgcaccc aaaagcacc ccaacagccc acaagtcagc caccatttgg tctccccaaa 240
 agctgatgcc taagttgcca attggccoct tattacaact tgaacttaac ctaactaaaa 300
 gccgctttta ttgattaacc caaaacatat ttttggtcag ccaactttac aaagattggg 360
 cccatatttt aaacaactaa caccttctaa aattgagaca acatgagtaa ttagatcctc 420
 tccatttggc cctaaaaaac tacaaccttg acttttctcc tagatactgg gctggattca 480
 aatagttgga caccg 495

<210> 30227
 <211> 392
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30227

agcttgattg ctttctttgg ctgaccctaa tatatgttac tcaaccatga aacaagtttt 60
 gttggataag catgggcctt gaaccaagta atttgatgca actatatcga gcanagagaa 120
 aagtcatgaa atactagagg ggtcatgcat atcctataat gacttgctgc ttggctctat 180

cagtaataga aacaatcctt ggatatagtg aagatagctg taaaccaagt ttgatttgat 240
 ccccaattta aaggatattc ttgcttgat gatataaaaa aagattgtga aggggtgatcc 300
 catggtcaga atgatgggtg catttaaagt tcttatgggg aatctattat ttatgtgctc 360
 ttgtgcaaca gggttgtccc ttgtattga gt 392

<210> 30228
 <211> 463
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30228

agggagggtga aactagtanc nctgcgacac anatacacia gcttatcaca tgtanctgtt 60
 ggcattgacat tngaaggga gaagcatttg tgttttggag tacttngggc cactatgtga 120
 tgccatggca aagtcttggg gtggccctg ccctcaactg gcattctctt ggcagcaaag 180
 tcaggtaatt gttggagaga tgtggtgact atgccctgaa ccctccactg tatgtcttga 240
 ctcatggcct cattcaaatt gtgacaacag gcccataag gtcggattca tgaacaccct 300
 ttataaaacc tagctggagt ttgtcattgt caatcacttt attactctat gaagagttaa 360
 aaatcagcgc ttctctatgt atcttcttag taagtttctt ttcttgcaat ttgcatagga 420
 cctctatttg tgggggcccg tctctccttt gcttatatgc ttg 463

<210> 30229
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30229

caattatgaa aattaatttt tgtgcgagat ttaatgttgt cacatgagct atacttattg 60
 gataaagtat ataagtatac tactaattac tcatacaaca tctaaattaa taaaaaagat 120
 tgcagtgtt atataataat tattagaaag atatatnaag agattaataa aaagatgtat 180
 taggttctat tgatagaggt atactaataa aaaaatacaa cgaaattcat tcagcatcgc 240
 tatttttttt ttaaatttag aagtatgaaa tgaaattaat ctcttttgca ttatacagta 300
 gaaatatata aaaaataaaa taattatttt atttatgatg gctcattcta gtgtatttca 360

cttaaagttt ctccattgaa atttctctta ttgattctgg atc

403

<210> 30230
<211> 489
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30230

gtgctgcgtc gtantctaca cnaaattatc taagctttga gatactangg atntgatgaa 60
gaaattgacc ttatctcatt ttttgaatga gggagcgaca atcggatgtg gatttttagta 120
tggtgatgtg agtcctttgc tacgtgggtg gggatcaatg atgaatgatt tttatgaatt 180
ccgacctcga gacattagcc attgattggt aatcatgtcg tcctcatacc aatgtgtgtg 240
gaggagaaat cctatattgg tgaatttcac ccttaggtcg ggagtatgta gggttactat 300
attctctttt aggtaatgtt acttgtcaac tgcattatat acttgcccta agcctttgta 360
gcaaaatgtg gcaaatgcac tgatattatg caagtcctat ggtacaatct ttaatattca 420
tcggacttga tgatttggtc ccttcccaat ttatgacgag gtttggcacc ttggcgagac 480
ttggtgaan 489

<210> 30231
<211> 474
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30231

aggctttctg tencctcgaca tncnannanc natnatagna nngnggacnc tccggaggcg 60
aaccgcgagg cagcgagccn ggcttttttt cttttttttc aagccaaaac tgagggggat 120
ggggccctat acctttgaca cactcaccga cacctaagtt ggtaaccaat tatggcacgg 180
ggtgaaaata actgggttca atctctatat cctatatttc ctccataacc tacgggggtg 240
aacatgaccc aggatttgga attgactttg tttgaaactt aatctaactt gcaaaatgtg 300
tctctatccg ctaagccttg gatggaaaac cctgcatctg gtattcaaatt attttctaaa 360
gataatttgt ttgcagtgga cctcaataat ttattacttg gactttaccc ctttatccga 420
acttttttat taaataaaac tcatactttt attaaattat gcactatata gccg 474

<210> 30232
 <211> 591
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30232

cagccccgat ggacatttgc ccagggttaen ngncnacnna tcagngcngn aannnanatg 60
 ctatggnnac cgcggcngga ggcgcactcn ctagtacgag ccctggacac gcgataggcc 120
 antgccanan gcanttgnaa ttttntnatg ntatancagn nacaccacgc cctnncgngc 180
 gggggacnac gtggtagttg taatactact actcctctaa ttaattgaac attccttgag 240
 ttcgattcaa tttagaaata aaaatctacc aaatagagaa atgagatcta tatatttaac 300
 tatacttttc agaaaataca tgcactctaa taggcaccaa agactatatg ctataccact 360
 cctaaatcta caattaaagc tacgtagaga agctaataaa aaaactttat attcaataga 420
 atgcgaatct tacattaaat aatcatacta atggatgaca attatacatg tgtcattata 480
 taagatctta cgaatttaaa atcacctcaa tatatatccc gagaagtcac atctacaata 540
 tcccggttta ttaaagttac ctattggggc tgattatacc cattctatcc g 591

<210> 30233
 <211> 395
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30233

agctttatct actttgatgg aatgaatcca tatggcaatn taagcactta acacaattca 60
 tggccaattc tactagtaat ntacaaatct tccttccttg gttgtgcatg cagtgaaaat 120
 acatgatgtt gtcgatgatg atatcangcc caagacagcc aggaaatgac attgatgttt 180
 atctaagtcc gttgattgaa ggcctgagaa agctgtggga cgaggggggtt ctagtgtntg 240
 atgggtttca gaatgagact tttctaagtc atgcaatgct gttttgtaca attaagtact 300
 ttccagcata taggaatttg agcagttaca gtgttaaggg tcatcatgca tgccccatct 360
 gtgaagaaga cacaagctac atacaactga nacat 395

<210> 30234
 <211> 422
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30234

agcttcatta agtttcaaga tngattcaaa gagtgggtgag gatatcaaag aagatgacta 60
 ancgctcata agtcaggaac acttcatgat aacacagctg atgatctcaa gaatcaaaga 120
 atgagtttaa gattgaatca tgtacacttc aaggatcaag aggaaagttg aattcaagaa 180
 tcaagtttca agattcaagt tccaagaatc aagatcaaga ttcattgactc acgattcagg 240
 aattaagaga agactcaatc gagataagtt ttaaaaagtt gtttttaaaa aataaactct 300
 gaatagcaca tgaatgtttc tcaaaacctt ttaccaaaga gtttttactc tctggaaatt 360
 gattaccaga ttattgtaat cgattaccag tagtaaaatg attctcaaag aacattcaaa 420
 ct 422

<210> 30235
 <211> 115
 <212> DNA
 <213> Glycine max

 <400> 30235

atgctctatg tgcgcactgt gctatcaata ctaaattcta gtagtgcctt tgccctcgatc 60
 acgcacatgc gtgctaagtt aggagcattc aacattgggg aatagtttga tcctt 115

<210> 30236
 <211> 358
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30236

agcttgactc gctcatatta ncatgannat ctactatgcg agtaatttat tatctatttt 60
 ccatctgcca acactcatga atagatatcc ttacgacccc actaatcctt tatatgtttg 120
 acttggaag caatttggaa cgccatttcc agtttgggtcg aaagattgaa gacttgatgt 180
 tcaatgatgc acctgaagag ggttggtgtgt ataattgttt ataattcctt atagtttctt 240
 gctatggcac atctagtcac gtctagaatt ctatttatga aacaatctgg ttgcaatgtg 300

tattagttac tctggttgaa atcacgaggt tctagatata gatggcggaa gagagata 358

<210> 30237
 <211> 422
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30237

agctttttta taattatcat cctctgaaat cagntcagag tatgcaagct gccaatacca 60
 tcacccaaaac taccactcac cacaatcaca tagtgtatac aaaaaaatta tagaatataa 120
 ccgctatcaa tctttcccaa tgtgttacta gataaaatta ttagcatggt tagattacac 180
 gcaagaatca attctaccct ataaaataat ggtgatacca tggaaaagta taagcaacta 240
 tttgtgggtt tgccttcacc atgaaaaaag tagctgttcc tagtaaagga cagtaggata 300
 acattaacat cagaaaggac caaagtcatt agcataggac caatattagc atcaagtttc 360
 cccttgtatt tcatacacac agagagctgt aggtcttatt tggtcgcgcac caccctttgg 420
 ga 422

<210> 30238
 <211> 459
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30238

gctataacac tcactcaact attagtatga aagaaataan tcttgctgat atncgaaat 60
 actcatgatt gtgtatgtga ctntaaaagg tcatgtgtgt gtcagtcact ttaaaagggt 120
 atatattttt ttttatttta atgtggatca ttcagataat agacacatgc accaagcatg 180
 aacgaaacta gaaaaatatg ttaagggggc aaaattttta cacattatan acaagattaa 240
 aaataactaa attttaatta tttattatct aaaatgtagt ttaataaata tgaaatatta 300
 aataacatat aaaagtggct atnattacct ttaatgcaag attatacgga aattgttgaa 360
 atttgtggta taggcacatg gtggaataga tcaaaaccat tgtttttctc taaaatgtgt 420
 gtttgggttct acgatggaga attatttcca atttatatt 459

<210> 30239
 <211> 381
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30239

tcttctttnc ttaagtggta tccggcatta cattgagact cgatccattg tcgataaaca 60
 cctttgcgac aacatgggtcc atacactgta ccgacacatg aagagccttg ttgtgtcttc 120
 tccccctctac ggggaatctct tcttccacag acgcgatata attgatgggtg gttatatgat 180
 taatgatgcc ttcaaaaccc tccattgaga tatcgtgcgc tacatgggca tcattgagga 240
 cgtttatcaa cagcgtacga tgaggctcgg agtttatgag cagttcaggc aacgacatcc 300
 ttgctggagt tttattcagt tgctcgacta ccttaaactc gctgtgttgg atgacgcgaa 360
 agaactcatg ggctcttcc a 381

<210> 30240
 <211> 528
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30240

cgggccctcc ccnnnnnnnn ngngttggaa agtcangnan cgnncnctn gatnaatnng 60
 agcgnagctc ccgtggagcc tcttgagtcg atgtgcacgc atgtttgttt cattaanagg 120
 cgtctcgac actcgggagg tggtgattaa gatcacaacg gccaaatcat ggccgctcgt 180
 atagtgaaga tgcatacctt atagcgagat gattctgcgg taatcgaaga ctcgtcatca 240
 tcctatcgca gctccttctt gatactaatt ctaagagcat cacatagaaa gcttctccat 300
 aatcatatct gagagtctt tgacaagcga ttcaggaag ctattttgcg atgctagagc 360
 cttatcgatc ctcacacctc tatcaagtat atgaactacc gctggaatta ttctcgaaa 420
 tgaataacga caccatgtat ctaccgtcct acatcatcac gtatgcaata ctatctgtat 480
 attctcgcgg tgtacatcgc acacacactc tccgcatact gtggaccg 528

<210> 30241
 <211> 375
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30241

attgtgcaag caatcaatga agcaaaacac accaaaagat tatgatgatg gatgactcaa 60
 atgctcacia atgtgaactt atcagtgttc aagtgagcgt ttcaatctat catgacatgt 120
 agaggcaaaa caaagatttc agatcgcaga atgtcatgag actattatct ccagaacaat 180
 taccatttc ttgagcatat gctacagttc agagaaaaat atgcatagtt gtacatacaa 240
 acanaattga cctaaaatat taaactagag acccaacaga actaacaat ttaacacgaa 300
 cgaaactatc agaactagca aaacgcaaac caatgacact cccccccccc ccatacttaa 360
 tacacatggc ctaat 375

<210> 30242
 <211> 402
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30242

tttcatcaat tcacaaacaa atctttgagt gatgcacttg aaatatttag aggattgttg 60
 aganagatgc ctactcatgg tttttttgaa ccaatacaac tcaacatatt tatagatgcy 120
 ttaagaccgc aatctaagta gcttttagat gcttcagctg gnggtaagat caaaatgaag 180
 acccctgagg aagcaatgaa nttaattgaa aacatgggtg ctagtcatga tgccattntg 240
 agagaccgag cccacatctc aacaaaaaat agtttattgg agcttacatc acaagacgct 300
 ttgttggcac anaacaagtt gttatctaag caactggagg cactaacaga anaacttagt 360
 aagttgcaac tcagcttcat tttgcacaaa cttcacattt tt 402

<210> 30243
 <211> 530
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30243

caggtagtnc cttgaaantc gtcgacacna nacatatact cnagcttcta tatcagctga 60
 agccgtggta tcaataagcg acaagttgag tcttattcan attatgagag ngatatctcg 120

ttatcttaag tgagaggaga ttctcccgag atatcttgag tgattgcaag aacacccttg 180
 gctgtatgca aggactttca caacctttgt gagttgccct cacttggaag agtgattgtt 240
 ttctctgctt tcgatcatca cgccttggtc tttcagacca caattccaga aaatccacct 300
 cttgccagaa ttatctcggg gccataactc ccattttacg cactcaaatt aagtgattct 360
 tgagcctaga ttgaatttca gaacgagacc tttcacctgg gtgtaggaat cacctcattt 420
 ggagccctgt agctgcaggt attgccattt ctatatttct gtgcagccac cacttaacct 480
 acggtgtacc atcccattca tgcattgtat gccagaacc accttattan 530

<210> 30244
 <211> 406
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30244

tgtatctttn tttcttgctg cccaatgcat ctagtagtgc ctatttcagc tgaaatggaa 60
 aagcaaccgc ctacttggga aaatgtatat tggtcccgat gaagacgacn gagacactgt 120
 ctctgtctgtg tcagaacttc cttgctgtgg catcctcaaa gactgtctcc gtctgtcaca 180
 ctcgactcac accacaccca attgtgataa acgcgctgct gtaatataat tanggggtgct 240
 ntaatattn ttttattaat atgattgaac caacaactca catatactac ctaactgaga 300
 gaattttgat tntgaatntg aatttcaaca cataattagt tgaattttta ttagataata 360
 tattattaga tctgtttatg ttagttcact attgatcggg taatac 406

<210> 30245
 <211> 507
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30245

aggagtgcac cncctangat tngctgcaca nanaatacac aagcattcgt atgacctta 60
 ccatgattnt gtccttgtag ctcccttaac caatttctta agaatatggg caacaaccac 120
 tttggtaac ttataatttg atgaccgggt agaatcaacg ccattacatt ttcttctta 180
 ttgggttttt ttttaacccc caccatcct atattatctc cagggtgtaa ttcttgtgt 240

ttctgttggc ggagtcttag gccgagttct tgcttgttct cgaggggttg gtttatttgc 300
aactggaaga cataggtgcg ggagacaaag acgtacgtgt acggatatcg ggctgggtctc 360
tggattggat tccattggac agagcactac tattatcagg tttacttgtg atgatattaa 420
gaaaagcacc aagaatctct ccagaaatac atagttggaa gaggtggtac aggattggta 480
caaggctttg cttctaataga agcgaag 507

<210> 30246
<211> 373
<212> DNA
<213> Glycine max

<400> 30246

cagccttctt ttttgattag gaaaatttta tccaccctaa ttaagttgga aatttaggtg 60
acagaaccaa aattaaacct gctcttaaata aaagaattat ttcttagaag ttaatggtaa 120
tggactaata attttaacct tatcattcca ttaccaatca ttattgatta taattttaag 180
aatatttcat aaaaatcatc aaatttatta tatatgatgg gttctgatcg gatgaaagtg 240
taaattattt tacagtgata atgtataatt ggttttctca atttttcttt gataggacta 300
gtgctgggat ttatctaaag aatgaaaaga tagtcatgta acttccaacc ctgagcaact 360
gaatgcaact act 373

<210> 30247
<211> 533
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30247

gggattcttg ccttgatcat gcgcacaaca ctaacaagct ttgccacaat cacacttgtg 60
ggtctgggtc tttcctcgta ttattattat caaatttgcn cccggcccaa aaccacttac 120
taccaggatt ctgtgctttg tctngtgaga agtagttggg tgtaccaga ccaagatacc 180
gcattcttgg gttgagaaca aagaattgcc ttgtgcttgc tgaggtgggtt attattaccn 240
cganggaatc atttgattgg ggttggcgct tattaaacat ccccaaaca cctgtcttgg 300
attgagatgt tctttgtacc aaatacccac attaacgtta ttgggaataa cctggtgatg 360
tgccattcat ttcttctatt ttctaaacce tttttgcacc atgttaatta ttgattgatc 420

ttaattgtca atttattacg caggtatatt atttgggccc attaagctta tgtgatgttc 480

ttatctatatt cagcattaat gaacattggc ttgatctgct ttggcttgat ttn 533

<210> 30248

<211> 517

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30248

ccccaggggg nnnaaggggt ggcttgccct gtatatctgc ganaccactn ctacngagc 60

tgctntgagn agacctagat gatggcagcc tcctattatt gtggcagggc ggcctccctt 120

cactttcttg tctccaacgc gacctctgac cactgttctt ccttcccgcg atgcttcttt 180

catggtccgc ctaatgggct tatagcccta aacatacttt ccacgaattc cctgggggtt 240

tatcaagcta gntatgctgc attgtctttt gctaaacca tcccgggtca taaaccgtcc 300

ctacataact cgggccatca taccgccgca tcggacagac aagggtgccc aaagaggag 360

tccacggagg aaatgctgac cacctcaaaa gactgganag cggtttctaa cgattcttct 420

gcggcttcca cataaggcat ggaggatggg cagcttacca agatatcttc ctgcctgac 480

acgataacca agtgcccctc cactacgaat ntcagcn 517

<210> 30249

<211> 368

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30249

caaacattta acttgaacat ctaaaaaaat attagcattt aagcattgcc cggctataaa 60

atcccaattt aaggatgtca cctaacattg atgaacttga aataccatag tgtgctgtta 120

tagattttga acttgtaaag ggggaaagca tacctacacg agtatgcttt tcctgtttct 180

ggcaagttaa gatgttacca aaacttaa attggttccat ttgacacaat atttaaatta 240

tccttatttc acttaaaata aacctcttta ctggttgatc tattttataa acctcanaca 300

tgatgcattg ttattggagt atgattgatc cagctgataa tttcccccat tgatggtaca 360

tattatat 368

<210> 30250
 <211> 393
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30250

gagcttcact tttttaattt taaantataa aatttcacatc gagttataag agtttttaaat 60
 aagatatgaa aataaatatg gccaatatg tggttttttt ggaaaattat tatacacatt 120
 aattaagtcc gtgccttata ataaaaccgg ggtaatatta tccgaatggc tactttttatt 180
 ctattctgtg acatgtaata ggttttgcac tcattacctc agggacgaag gaattaagat 240
 gatttttttg cttcattacc tcagagacca ggattagggg tgaatattgt acggacatag 300
 acgctcatga tctttntatc ttaaaagaaa tatctctgcg tgctttgaag ataacaatat 360
 agactctatg aaaacatttg agctatactc gcn 393

<210> 30251
 <211> 489
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30251

agaaggatga ccatgatant cgtacataat attcagctga cacatccctt tgaaatgaaa 60
 agcaagaccc attattgtct ttatccctaa ccccgctgct ggattaacct ggagacacaa 120
 aaatttatga atattccgac cgattttttga attaaggcca accatgggtga aaccaaatgg 180
 ttaattggga gggaatattt gaccaattta aatatattat cttacccttg gagaacctat 240
 cattttgagg aagaaaaaat gggtacttca tggattaact tgctcttact ggtgccaatt 300
 aattatatta tctaacttaa ttattaagcc aaggatatat acttaataata gaatgcattt 360
 cccagtgggt taaatcattt caggtggctg aagaaagcat gaccaaccag ctttaaccgg 420
 catcttaatc cactttgcat ncagagcccg atttgaagat taatntgaat gggtagaaat 480
 ataataatn 489

<210> 30252
 <211> 249

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30252

acactanaga gaagcaatgg cgatattgta cacacacaca ttctctatct agaataaacc 60
 tanaatgtgc attttcactc tactaactta aaccctatgc aggggaataa aaagtaggtc 120
 tgatcttata tgccaacacc ggcagtgtac tagaatagat cacatccttc ctcacattgg 180
 tcttccataa agaaagcggg aacggggaaa agtaaagagg gatatgtgac ggtgctttgt 240
 ttctgtaaa 249

<210> 30253
 <211> 395
 <212> DNA
 <213> Glycine max
 <400> 30253

ggcggatgag aaaacattgt ctatattcca tctccaactc cagtaggcct cccaatcatt 60
 ccttactttt ataggaggaa tggtagggac aataacctca atgccgtttt gtctaggaac 120
 acacatcatt cctagtctc ttccttcttg attattatga tctctatact caattgaacc 180
 acctctcatg gagcgcacatca tctcgggtgat cattaacctc tccaaatgta gcatcaaagc 240
 ttgcatgaaa gattgcgaaa gcccactcc ctcattagga gtaataacctg gcatctcaaa 300
 caagcatatc aaaccttaca agacaaatat aggaactggg tgaataacctc acccactcga 360
 gaggatcaca caataatggg ctgtctctaa cgaac 395

<210> 30254
 <211> 513
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30254

aggaatgact tttctanatg attctgagaa ncatcatnna cgcattctna gactccanac 60
 actgcgngn atatcaaaga ggaagtttct tatcacgtct tngtatatga caatttacia 120
 gcaaactata gagacagtgc atcagacaac gatataacag cgaatgataa atgcctccat 180
 catataaatt caaaacacga ggggcatcaa actgtcatca gtaggaaaat gatggatgat 240

atttatgcat taggggcaaa atgtagggga agatgtggat tactcactag gacatcgtaa 300
 cttgaaagtt accatgggtg aggaaataac ataactgtca gattaaaaaa gggccggtca 360
 cacacaagga cctcataagc attcacaata gtatgactan attgaaagta attattcagc 420
 gttaccttca actattttctg ctggtacata agcagcacat tatcattctc aagaagaatc 480
 actccgatta caaattacaa aacaaatcaa tag 513

<210> 30255
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30255

agctctgctt atttggctct cgccagcgaa aggatcgaag tggatctgaa aagaggcaaa 60
 tctaatactc ctgcttagac gaatgagaaa actgnggcaa ataaagaggg tgaggatgag 120
 ggacaaaccc atgctgtgac tgccattcct atacggccaa gtttcccacc aaacccaaca 180
 atgtcattac tcagtcaata acaaaccacc tccttaccce ccaccagtt atccacaaag 240
 gccatcccta aatcaaccac aaagcctgtc taccgcactt ccaatgacga agaccacctt 300
 tagcacaac cataaaaaac accaaccaag aaatgaattn tgcagcaaaa agcctgtagg 360
 attcaccoca nattccggtg tcatatgcta acttgcctcc atatctactt gat 413

<210> 30256
 <211> 438
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30256

actaagctga gctcactggt gtgcccataa agtccacga aatttggtcg gccatgctct 60
 tccttgcgag cctcttgggt ttcttggtca aaggctcttg cggtagctgc attntcttct 120
 cgtaaccggg cacactctnt ccgaatgtct gtagcgacca acttgaatgn ttctttggca 180
 agtcttgcta ttcttagttc tggtttgaga gcttagactt cttcatcctc tttcggagct 240
 ntgaaattct cttcgttgat aatctttaac ttggagagcc aatctaacc tctgtgaaga 300
 actttcagcc attcatgata accaccgatg aagccattac gaatgcccct aagttcttta 360

tctttcctta acgagctttc ccacgcctta tggactcttt gtataacctt gaaactttgc 420
 gcgccgaaat ctctcaca 438

<210> 30257
 <211> 366
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30257

gtgggtacct atnttgaatc tgcgatgctg tctctacata catanaacag tcccaccatc 60
 ccaattgtgc aaaaccatat tcatatatca ttgcggcatt tcaccgagca cttggtgggc 120
 gcacgtttgg acataaatcg caagagaatg ggggcaatgt ggcattgcctc attgcttcag 180
 aacacaacat aggcctaagg ccttctcatt caaatcctca actcaagaca tcaagcatac 240
 aaacaaccca caactgcctc accaatgtaa gcatgttctc acaattagag caccagaaga 300
 tgaagaatat actccaatgg gaagcataaa actcaaggat ngaatactta cttgttggag 360
 tgagta 366

<210> 30258
 <211> 492
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30258

aggaggatga tnncttagac tctganncaa attacctact ccaccgagac gctnnaagta 60
 gagctggctc atattcctgt ctnnnctga gcgcacctcg ttggatgaga actagagcta 120
 tctaccaccg gctataatag ctaagctcac ccccatgaca aagaagctga aaatgacaaa 180
 aaaaaaaaaag tacgttatac acaataactg agattgcgcc gaattacaag gcgtaaacc 240
 tatacttact aaatggcgca aatacaaggt ctagacgaag gaataaccta tgtaaatatt 300
 tacgaagata agcgggctca tactaagccc atgggctgga aatctaccct aaggctcatg 360
 agaaccctag ggcctttcnc tggatctcta gccagctca cttggagtct tctaaccgat 420
 gctcttgag ggtaggatag catcattccc tccaccttag gaaggatgtg acctaaatcc 480
 cgagttcatg ag 492

<210> 30259
 <211> 524
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30259

agacgttctg annctgagta gnatncctag ggcgnctact ctgaccnngc catactatcg 60
 agnngagccg acaggcaggc aggcaaaacta ttttgncttg acnagacccc nnagcttagg 120
 agagatcagc tctacaaaaa taacaaccga ggagcggaaa gtataaaata ttaaaaaacta 180
 atataaacga tgatgntaat gtaacaagtg acttcgaata aacatcggag ggaaataata 240
 ttactgctag gctacatact tatattgtac agagaactac tacaagtaac cttacaaaac 300
 gtgacaacta tgtagaaacg actaaaaaag attatttatg caacaatgag tacaacttta 360
 cagagataaa atatagttga aaatataatc gagcttaatc tctctaattg gatagtaaga 420
 caaatgctca tatgacatct ctatcattta taacgtgcc aatttgagc ctgggtattg 480
 ctcatatgtg cagtactttt tacagagcat gtccagccac gccn 524

<210> 30260
 <211> 506
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30260

agagtaatgg ctnatgagac tntcggacac ataatctacg ctagcaagaa ccangcggga 60
 nggagaatga agaggcgaca aaaatgancg atgtgcncn cgagcccaga agaggtgatt 120
 gagcctggag accaagacac ctatgaattc ctacaccgat atcaagatgt tgtccggcta 180
 caccaacgac tatggcatat cagcaaggat gtacatttct tcctagtcat acgcccggcg 240
 catggatgac ccactaagga cttctgcca aacatgatta tgattctccg cgaatcaact 300
 tcacgcatgc agcctagccc tccaagcact gagaccagac gaagcccga aggacaacca 360
 catatcctac tcgcgaggtc tgtgctacca attctttatt gctgcacgag aacgaccact 420
 cctttccact caatgaacac aaatgaactt ttctctgct accgtctctg agtcagatct 480
 ataatcactg cacatatcca tttagn 506

<210> 30261
 <211> 415
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30261

agctnttctc cctcattctc acattgcttt ntctcccttt ctctccacc attgaagcct 60
 ccattanagc tccaaacttt gctcaccatt tctgctccaa atcgcaaaag gaagctattn 120
 tcggagtcgt gaagcgcacc tctacgttgt gggaacttca aatttaggtt tgggtagact 180
 tcttctcaca taaattntcg tgggtattgg gttttgggag atatgatggg tagttgtact 240
 aagtttatgc cttaaggtag ttatttgtga aggaatttgt tgaaagcatg ctaaaattat 300
 catgtttgat gtgagctaaa tataccatt ctgttttaag gttntataat gatactttgt 360
 gatgcttggtg tgctgaaatc gttggtagaa aattgataga gatggagggt agagt 415

<210> 30262
 <211> 210
 <212> DNA
 <213> Glycine max

<400> 30262

cttttgagct agaatgtgat gcctctggag tggaattgga gctgtttgtt acaagctggc 60
 accctattgt tatttattga aaacttatat tgcaccctaa ctaccacctta tgataagagc 120
 ttatgcctta taagagccct ccaacttggg acataccttg ttccaggaat tgcattctag 180
 tgacatcatc acttagtcat tgatagcaag 210

<210> 30263
 <211> 316
 <212> DNA
 <213> Glycine max

<400> 30263

tctgctgcc aacatacaacc tttgcccttg catgcaacta cctggagcaa gtgagcagcc 60
 tgaggcttat gctgcgaata tatacaatag acgctgctca agccgcagca gcagaatcta 120
 ccacagcaga acagttgtga cctctgcagc aacagatata gccctgcatg gaggaatcac 180

gctaacctca tatggtccag cccttagcaa caacgacaac agcctgctcc ttacttccaa 240
aatgctgctg gcccagacat accatacatt cctccaccaa tccaacaaca gcagcaaccc 300
cagaaacaac caacag 316

<210> 30264
<211> 379
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30264

tatggtgttc aaggtggatg ntgaaaaagc ctatgactca ctctcatggg ttnttttggg 60
ttatatgctg canagaatgg gtnttttgcca cacatggaga cactggatgt ctgcctgtct 120
caagtcagca agcattgcta ttcttatcaa tggcagtcct acaaaggaat ttgctcctac 180
tanaggtttg aggcaaggatg atcctttagc ccccttactc tctaatatag ttggagaagg 240
catcacatga ttgatgaagg aagcagtcaa aagaacttat atagaagcta tatggctgga 300
aagaaaaacg aaccatttaa tatcttgagc tatgcggatg acagcaattt tgtgggtgag 360
gctgagtggg agaattgta 379

<210> 30265
<211> 173
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30265

gacctggact ggaaacgatt atcatgtact acagcatgat gatggaaagg atgacatcta 60
ttcgcttatt atcactgagt gaggcataga gaccacacgt ttatggctac agaagatgtc 120
ctaataaatt atccacgtct gccatcatca agtactgttg taatgatcag aan 173

<210> 30266
<211> 275
<212> DNA
<213> Glycine max

<400> 30266

ttgaaaccac tttctcactg cgttgaactt cctaattaaa tgaaataatt tccctataat 60

taccatggac aaattccaat tgtaaagatc caattcttat ttacctaaaa tgattaatga 120
 ttcactaaga catcatcttc tcgctgcttt tgacaatgag tatgggtgaa cgaagccgta 180
 cactaatcca atacacattt aaaatacagt atctacgaag tgatcttacg ttgtctccaa 240
 cgagcaatgt caaccaaag ttcataacac atagt 275

<210> 30267
 <211> 478
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30267

aggaatacta nntc gattn tctc acanan tate nnanat aaggcatccn nggccggnc 60
 tangaggaat cngagacat catntattnt tcnnngaanc ccanncactg gaggggaccg 120
 aacgcaggaa tcaaaccgac cgtgataaca tggaatccgc atattttatt gtacaatgaa 180
 atatggaacc cacctctggg tttcatattg gtgacccatg cctcataaca tatgagccat 240
 cagtttagta agttgaaaat attgggcaag atgtgttggt gtgttgagcc acgtgatgtg 300
 aacaactgaa tgtataccat aatgattaat gcatggctat ggagtttaat tttatattgg 360
 actaatattt tatgggacat actactgata aaatgtgac tagacatcat tgatcatgca 420
 agatcctaac ctttaacaca gtttggaag attaagtatt tgctctatac aagatctg 478

<210> 30268
 <211> 389
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30268

agcttcatac tttctaatta atagcttcca aattttcatt ttgaagctat ctcatgagtt 60
 atggctcttg catgtctccc atgttttgta ttcttccttt atatctttag taatatatag 120
 acttgtagca taagatcatt acgaataggg tcaactaatat agttggtagt ctttttccta 180
 attaagtga tatattactc ttatactatt ttctacaaga ttctttctaa aagctatcat 240
 tttctattca tggctagaag acatggtttt atgatggtga tttggtgacg atnttataat 300
 aatcaacatc attaaggagg caatgacatt tttgtaaata ccagtcatat ttcaacgact 360

gtgttccata aaacgatgta gaaattgca

389

<210> 30269
<211> 250
<212> DNA
<213> Glycine max

<400> 30269

tgtagcattg ggtatctttt gtgatcgaca gcaccaccaa gaacacaaat agtgtcgaca 60
tgaaaaaac aggttgatgat ggtagaattt cttcttcttt gcaaacaaaa ccactatcat 120
agatcctctt cttattgacc agggtgagta ttttttttag tacgttcctt cctgctcctc 180
tgggttcatt tacctattta cattggatga gttttttatt cgggttttagt tatcactgca 240
tgttcattgc 250

<210> 30270
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30270

cggggaaggg gtnnnnttnn nnnnnnacct tgctagtcag gcataccgga gtanctgagg 60
gagcaacctg tattgtttgn gttatttgcc gcaagaaacc tgccattttt cttatctttc 120
ttcagggccc tatggtttgg cactccggcg cttacaatat gggatggttg ttccgacctt 180
tggtttaccg attcgcaaaa ttggaggatt ggtagtggc tcgttttctt cacccttcgc 240
aaggatattt tggccaccag tttctattct tctaattgta actgtgaata gtggtatgat 300
catgtggagg cctcttattt ctaattccaa actttggatt tttttttaac ctctagtttc 360
attgccccct aactggccgc tataccattt tcctcgacct ttttgtttaa gtgaagggtc 420
ttatggccca ttttctgcgg ttgatcccc t 451

<210> 30271
<211> 239
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30271

gtacaaaaaa aatcctgcac atattttcac ccttcaactct ataaatacat gaaatcgatt 60
 attctgacaa aatatatgcg tccgcgtggt cggtcgacaa actgtntgat ctgcagaact 120
 gcataccatt tgatatcatg tttgctcatc cttgcgtggt cctctacaaa aaaaaaaca 180
 aaaaggggga agcgtgaaac ttcatactac attcttagtt tcatgtgtta cgcaccacg 239

<210> 30272
 <211> 423
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30272

agcttggttca tagagatcta ggaagaacgc cgcgggccgca gggactagtg ccgctcccga 60
 gttcgatagc catcgtttca ggagcgctga gcaccagcag catttcaaag ccatcaaggg 120
 atgggtccttc caccgagaga gacgcgtcca gctcatggac gacgagtaca cagaatttca 180
 ggaggagata gctcgtcngc gttggatggt gctggtcattg cccatgggtca agtttgatcc 240
 cgatatagtt ctcgagtntt acgccaatgc ttggcctaca gaggagggcg tacgggacct 300
 ccggtcatgg gtaaggggccc agtggattcc ttctgatgca gacgccctca gtgtgacatc 360
 ctgaaaattt ctacctgaaa ttnttgaaac gatgtatttt gaatgattat atatatataa 420
 gta 423

<210> 30273
 <211> 447
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30273

agactctgca ggtagatttt agccttagtt tcactttagt ttgtagtcaa tncaattaag 60
 aaagagaaat gccaaagaga aacgtccgat tgattttttt tgctttattt tactaaaagg 120
 tattttttga ttatgatatt attattatac ctcttttttg atttccaacg tggttacagc 180
 acgaccgaac ggtcggattt cattataaca gaaattaacg gatattacag atcaaattgat 240
 ccgtgaaaat ttattttatt ttttgattag gcgagagatg acttaaataa atgactgaaa 300
 cacgtcaaaa gaggggtacgg gaagtaaattg atacaagata ttaaagtaca cgaatcagat 360

ggagaccacc acgaatacat aaaatgaatt gaagagctca gtttgggtac ttaccggttg 420
ataaccgatg aaaaacgaag aacgaac 447

<210> 30274
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30274

gccatgcaag cttgacttct aactcaaac atggcaaggt tcaacacact ggtcagacaa 60
atcttcttca ccaaataacc ctatcacaaa gcataanacc annataaaaac ctacccatca 120
tatnntccc aaagccccat acccācgaaa aatgtaggtg agaaagaagt ctacccaaac 180
ctgagatttc gaggtccac acgtagagat gcgcttcacg attccgaaaa tgccttcctt 240
tcgcgaattg gagcaaaaat ggtgaccaa gggtggagct ttaatggaga ggaagaagaa 300
agaagaagca acgtgaggga gagggagaaa gcttctgaaa ttntctgttg agtgaggaga 360
gagagaaaac agctnttttg tttaaagagg atnntctctt ttctattatt ntattntaag 420
ctatgccaca tgtctccatt tgagtgga 448

<210> 30275
<211> 434
<212> DNA
<213> Glycine max

<400> 30275

atgaagagtc caaagcaata cacatatatta acaaatacaa aggtgaatgt tttttaacac 60
atgcacgcaa acgaacataa aggccaaaac gaacacatgc atgcaaacat acataaaggc 120
caaacgaac cacatacaaa cgggtaaaaa aaaagaacaa aatagaaaca attgtaggca 180
tcaaaactga tgcaatccta ccccgcaagg gcattggata gaaaactcca agtagattga 240
gccagagatg caagagaagg ccctagggtt ctatgagcc ttaaggtaga tttcggggccc 300
atgggctaag tacgagcca cttatctatg taaatattag attaaggttt cattattctt 360
gggccttgta ttttaaggctc cataatagag gtagaggacc ctagaaatat aagagttttc 420
agcccttgta tttta 434

<210> 30276
 <211> 332
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30276

atcttgagtt gatgaagtgt tgaagggtga aacttcctgc ntttattggt gaccacagag 60
 tggtagctgn agatatgtnc gcggggtcat gagaccttgn ggacgtcang tgggggtgcta 120
 tttgccccaa ccaaacttga ccaatcccg cccacccgg gtgtcgcaac ctacccttcg 180
 gcgggagggc gacgcgtgac ttgctgggatg cgtgttccac ggaaggaata cgcgcgaggt 240
 cgccaccaac gtttatttga ggaaaacgtc ggaaaaaccg gaaaagacgc gatctacgaa 300
 ctttttagtg aaagggttcgg gagttgtatt ta 332

<210> 30277
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30277

agtcattggc gagtengact ntctgcgaca catttttact caagctgata cccgcagaga 60
 caacgtcgtg tttagccca atcaatcgn gcgacaaccc gnaacgcgcg ggatttcgta 120
 atctccgcct ctcaagatct gtatatggac tttagacacg cagatggcgg ataacgcgag 180
 tggtagctgt ataacttttg ctatctgtaa aacaaaacgc tgtagcacgc aaagacaacg 240
 gcggctttgc gccttcgcaa tgcgggtcgaa agcccgtgac accagagata tacatatctt 300
 tcgcgtcca agaactgaca tctgactttt ggtcgcgcta ccggccgaat acccaagggg 360
 atccgataaa cttgtgctgt ttagacgat agctggtaca cccaagacta cgtngggttg 420
 cgccttatca tggcgggcgac caccgggtgc ctcggg 456

<210> 30278
 <211> 500
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30278

cgccccgcan nncnaaggat gttgaaagct anagaagctn cgcaattatt cattgncccc 60
 cgattctctg agccgactgg cgcgtggcca gattttttct taattaatnn ccaatattta 120
 tgatggtgta ataataataa attttaagac catcctatta atatttatcc ggtggtnttt 180
 tatattaata tatttaagaa taaaatatta ttttggatat tccatattca ccggatcgga 240
 ataattgggt tttttatata aaaatccatc tcttaggcct attttaaatt ttcattaacc 300
 ggaattntta tttattgaaa tacctaaaat taatttaggg caacctaccg gcgcgtggat 360
 ttaatattat tatcaatcct attatatata aatggagtga ttccgaaggg acatgtataa 420
 aactgaaagg ctagtattgt gcaaaccagc gcagaaaaag gatcgtgaat atggtacaaa 480
 tattttgaga gaaacacaan 500

<210> 30279
 <211> 440
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30279

ctatgaatat attaagagga attgtatata attctggcga tnagaagttc tcccacccta 60
 caaaaacaat tntcatattc aaacagaatg ggtaaattat aaaaatgctc agattagctt 120
 caactatgca gacaaaaaaa aaaatagaag agtaagttcc tttaactgtt aaaatatagg 180
 acttggtgca natttaaata tcatgcaaaa aaagaagtgc agagtgcggg gaaaaaaca 240
 agatgtgatt ctgctttcaa ttgagaaatt gtatttagtt cccattaata aaatactact 300
 ggttcaaaaa actaagatta gaacagtgtc tgcttcttag ttcttacagg aggaatctga 360
 actacattaa gttatagact agagtcattt cagatcatcc ctttttcaac tacctcatag 420
 tccttcagct tcttgtctag 440

<210> 30280
 <211> 381
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30280

ctctatncct ccatcatcct attcactaag aggtagatga tctctattgg gcatttgtgt 60

aaaaaagctt accctaactt tgctttttaa tgtaaacttt gcattatttg atgaatcagt 120
 ttatagagat agaaataata atgcgttgaa tatgtaattt ttactatcat cgaatcagat 180
 cacanatagt gtgtatgata aacttggttaa atttatacat ctaccttaaa gttaagatat 240
 tatttcaaga ataatacaat gatataagag ttaattgcga caaatgagat ataaacttct 300
 taacacgtta gaataggact actactcaag tataccaata acattctgga tagttgataa 360
 tataaacttt accgctatac c 381

<210> 30281
 <211> 369
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30281

actacaaggt taatcaatta cccacctaaa aactacaact attgaattaa ttatctactt 60
 cgctgtcatg ggcttaactc gtggacaagt aaggaaattg gagtaaccag agaacaatcg 120
 agaaactcan gcaaattgtg cctaccacat acaccagtac atcgaagtac ttactttaag 180
 ttaattaacg taaacaaact ctgctgtttc tcaattctaa cccaacaacc ataccaata 240
 aagagaacaa ctgctggatt agctattttc attaattctta ttctatatat tattgcgaag 300
 ctatctgata tatatcctgg acgaagagta ttccaagnct ttgaggttc tatataattt 360
 ttttaaaat 369

<210> 30282
 <211> 395
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30282

ctgctgcttt ctcangettga gtatctaaat atactaagga gaagatccaa aacatcaaag 60
 atgtacaata gaaggaacga ataacaactt attttaaact catctatggt gacctatggg 120
 acatggtgga aaatggaaat tacattccat ataacgatca gttaaacaaa attcctaaaa 180
 gtcaatggac aaaggagcaa tctgattttc tcaactcaga gactccaaat gtgatgctat 240
 atgctctatc agaagatgag tacaccaagg tacacaactn taaaagtgtc aaacaaatgt 300

gggacactct agctgtaacg tatgaaggaa cgtgacgggt aaagaagaac aaactaagtc 360
tgctcactca taagtatgaa atcttctcta tggaa 395

<210> 30283
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30283

cgccacggaa aaaaaggaag ggagaatddd cangtananc acataannaa agcnggacac 60
gagatccttg gaggggacct gccgcatgct tctttgtacc tctntngcac tgcacactga 120
ctattcactt ttacttatgt tcatcaatca ccctaacaca ttagctatga gaataattta 180
tcaagaaacc ttttcatgtg gccatttcta atagatcgag gactcttgag tacatgtgaa 240
ggctgctata cagaagtgga acaattcaat tatagtatca ttttactacc ttacactcta 300
agtgcgacag atactctgtc catagtgact ttcattctct cataagatgc aaagagtgat 360
atgtaccgtt acaaaggcat ctgttatgct tggatagcta ctgcacagtg gtattggctc 420
tataacctaa agctctgg 438

<210> 30284
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30284

aatcattcaa cgcattcaga atatgtgttc atccttanta aagctactct aaaataaatg 60
acacagtcgn cagcanagag taagtatcag atgatagggtg caccctata gatnttaatg 120
ccataagtat gtccttgccc ctccaacttc ttaataagag cttanatccc ttcagaacaa 180
aggatgaaca aaaaatgaga gatgggatct cctagtctga gacctttccc ctggataata 240
ggaccaacca agctttcatt gataataaca gagtagacag attggatgag aattaaaatc 300
cacttaaccc aagtcgcact aaatctcatt ttggccatga cgttttttaa ataattccca 360
tcgacttggc catagccttt gttgatatcc atcttcagcg caacttgtaa cctcccat 420
gttacccttg accttacact gcatatgatg gagaat 456

<210> 30285
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30285

agctatgtgt ggttcttcaa tggatgaatga gggaggaaga aaagcaacgt gagggagagg 60
 gagagagagc ttctgaaaat gtggggctga gtgaggagag agaggggtgc tttttggttt 120
 aaataaaagg gttntctctt tttctattat tntatttaag caatgccaca tgtctccatt 180
 tgagtggagc aagaagggcc cactttctct ttttgactgt gacccatatt cagtcacaaa 240
 agtgagaaaa atctgacctt tgaaacgcta aaatcctgcc tcggtttgcg tgccgtttct 300
 ttgattccag tttctcgcgt ttctctgcgt ccgccggggc cagttttcga aagcaagcaa 360
 tatatatatc anaacgctca gaatanaacc ccgaacgtgg ttcagagggt ggtttcgtta 420
 a 421

<210> 30286
 <211> 423
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30286

aggagtttat tctgcattcg gcannnctat ttatcngncg aactcagagg ggnnnngaag 60
 catgcaagca agcagtgata tttcnnnncc ccgcnnnagg aggggggttg tactatcatt 120
 ccctaaaaac atcaacatat caacgttact ctttatattac atcatgactg ctgacgaagt 180
 ttttagctgc actctgagat attggtgacc tctactgctag agtgacgacc tgtcttatgc 240
 tctccaggct attcaaaatt tgcttgtctt tcttgcgacg tacttggtta tttcttccat 300
 caatgactca tgcctgaag tgtacatagg aatataacgt gttatgaact ctatttattt 360
 gaaatttacc ttataaaccg actaaggtaa caccttgcg atgaaccctc cttaagagtt 420
 cag 423

<210> 30287
 <211> 337
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30287

aggagattta tctgactcan ncctatnttc gacaacaacc gncgggggag gaggatttta 60
ttttnnnnnnn caaagggggg gtattacatc accaaacacc agcangacac aacatgggca 120
atccccaatg acattgtggt ggcaacacta caagtaatat actttaatga cttgagattc 180
ttactgtaga gatctgattc aatacaatgt agacctttcg caacagacca tacttgacta 240
ccatatcaaa aacatcaatt tctcaccaat taccaacttt aactggatta cgacttacat 300
gaagtacccc acatgcctgt cctttacaac agtcctc 337

<210> 30288

<211> 409

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30288

ggacctgcag gcaggcaagc ttgattttctt ttctcacata ctggaatcga ttaccagaga 60
agtgtttcag aaatattctc acagtccatc ttttacttga ttctgatggc tgcaaagcct 120
atattatggg aactggacac aaantgccaa gagtctttca aaaccaaag gtattatcct 180
ctaaaaagca catcgtttta tctctttaac aaattccttg gccaaattac ttgtgattca 240
ataaggaatt atttgagtgc tcaaattgtg caatctatct ctttcaagag agatttcttc 300
ttttcttctt cttcattctg aaaaaaggga ttaagagacc gacggtctct tgttgtgaaa 360
gaattctaaa cacaaaggaa gggttgctct tgtgtgtcta gaacttgta 409

<210> 30289

<211> 512

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30289

aggttggctc cgctgagant ctaccanact actcnaanct angacactcg nngagagggn 60
cttacagggtg agagaggagc aaataatn tctaacncnc cccgaaacaa cggcgcccct 120
ggcagatgat cgcacacgcg aggccaggaa ccccgagatg atccgctaac actcttgctc 180

gtgagagcag aaatgacaac cagtgggtgga caagaangtg agattccttt gtggagccgg 240
cgaactgcat gatgaccgtg agattatttg ggagagagtg tgttttgtaa tcaactgctg 300
cctagcaggt ccggaattct ttttggtgat ttggagactg aaatcacata tttaatcata 360
tgtgtgaaca aagttattcg tcattatgtg aatgatgtgg actacngac tatatatata 420
tgtatatata tctcgtatgt gtgtatgggt ggattccctc aagcataggt gcactgtcct 480
ggggatgtat atcggtaaaa cgattcgttc at 512

<210> 30290
<211> 349
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30290

tccttcctta acctttctag ctgtgcattg gtgtattttg atctcctttt ggtcctctaa 60
ttgtggaatg tgttcaatat gtgggcaatt tttgggttgt ttccttgctt gattgggtta 120
gaaattgggg gggtttgtat ggagatgggc cctangcctt ataatgcatt ttttgaagca 180
atgagacatg ccacatttgt ccccggttctc ttgctattga tgcctaaaca cgcgccacc 240
aagtgttcng tgaaatgcc ccatggcatt agcgcgtgggt ttttgatgga aacaacccat 300
ggagcatttt ggtttgaca tatnttccat tttttgggac atgcattca 349

<210> 30291
<211> 366
<212> DNA
<213> Glycine max
<400> 30291

acctttgtca atgatattct tcatgcctct taagtgcaga agtccaaatc tttgatgcca 60
tattttgact tcatctttct ttgcagggtg gacatgtgga ggagtaactg gttctttgag 120
gtgtccataa gtagcagttg tcccttgatc tgetgccctt cataaaaaact cattcttctc 180
attggcacca agcattctga ctttgtgaag tttacattga atccttcac acccaactga 240
ctgatgctga tcatagttgc agtcagtccc ttcaccagca gtactttgtt cagactagga 300
agccatcatg gactagcttt cccattccag agatctgtcc tttagagcca tctccaaatg 360

<210> 30292
 <211> 501
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30292

aaggagttca tcttcttgca tnacgacann nctattncta tcnngcgaca catncacang 60
 gnngacctgc atgcgtgcaa gctttcgttc ttctntnnnc actgctgcga gaaaggggggt 120
 tatttatcca agtgagatta caagccccta acaactgtgc ttgacaacac gcctaagtcc 180
 gacacagatc aggtgcttga cgatgtgtat ctgatagaga acggcacagg ttttttcaca 240
 cggatgttga acttaaactct gtttacacaa acatcctatt tatgactata gaaagtgaac 300
 aacctgcctt gattagctgc ctgctctccg accaccgata tgaagtagat tgcgcttact 360
 gtgcttctcg tacctgcaca ccgccacact tctagttaag acaactctcg tgcgggaaac 420
 tngatgcttg taaaagtcta ccatatcagc ttaaaaagga gaactacttt gcattgcaga 480
 tgggtctaacc atctatcacc c 501

<210> 30293
 <211> 412
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30293

agcttctatt actntattga cacacaaaat acctaatttg aatgaagcat ttgatattatt 60
 tcttangatg tagtcttaag atgcgaggat gaaatctaaa attagggttaa taaaatttgg 120
 tcactttttg aaataatatt gattgaagat atggatgaaa ttgaatattt aatattaaaa 180
 aatttgagta atttaaaaaa cttatataat tcttttataa ttataataaa agtgggtaca 240
 taagtaaatt attcttatga tgatcaaag aatcctataa gtatatgtaa aacctacaaa 300
 aattattctt tccacaattt acccatgcat tgcgcggaaa aaattgacca tagttttttt 360
 ttttaactta aaaaattgac catagctgan atgtaatcta gtttcgctta ta 412

<210> 30294

<211> 259
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30294

 ggagatcagt ttatcnanct tttcggcacg cggcgggggg gagctttttt cacacgcagg 60
 ggttgtagat accccccctg cacactcctg tgcggctata gcgtggacaa ccaggtttaa 120
 taagttatct acaaggcgca tcttgaaggg caatatgatg aagaaagggg taactgtaga 180
 attataaaaa ttatatatta tctaaggaaa aaagattgtg aaagaccagg gggcctggta 240
 atcaagcaaa aacgacgac 259

<210> 30295
 <211> 388
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30295

 agatcgtgac gttcttgenn ncntatnnat gnnccacnt ctgagcgccg gggagggctt 60
 tttcattttt tncnacncag ctgggggtgc gcagctggag aaaagaacca acaagaaaca 120
 gcccatgatg cgggggtggtc tttactgcag agtggggaac aagaaaaacg tttgactgcc 180
 tttggaagca ataactcacc ccattccatg actttcttta agtggagttg ccgcgcgcag 240
 gtgggtgaag cctcgagagc agaagcacca agaggaagag agaacaccgc acgacctcta 300
 gaattggata aaacaatcta cagggtgcta aagacgctac atgggcttca aacatctgct 360
 tccgattaaa tgggtgcatt actctcgc 388

<210> 30296
 <211> 372
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30296

 agcatatfff tcattaaaat agataaaata ctatgttcag atatacatgt ctacttgtga 60
 agagatagct agctatcaca tttagctatg gtgatcagct tcataaagag tccttctaaa 120
 cccaatcaaa gcaacaacaa agtaacaaat ntacagaata gatcaagtga aaacacttga 180

agcttaaaat cagtaatcaa tatgattgga tagaactata gtcttatcta aatcacangg 240
 caaaccacaa cttgcaatan aggcaaagta actaaatagt gactactata gataacacta 300
 atcaatttcc aaagtgcata caaaatatat ccaaattgtgt gaataataat agtcatgata 360
 atgaacaata tg 372

<210> 30297
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30297

atccaagctc atcttggagg agaagctccc tcaaattggc ttattcccta gnggaagaca 60
 cctgcggtca cctattctcc tttgtcttcc gctgcatctc catggtggaa aatcaccatt 120
 aaaggaccta attgaagctc aaagatccag tctccataga agccacacaa gcaagcttcc 180
 atcanaatta gacctacgta gttctttcat aagaacagaa cgttgggttaa gttgttttga 240
 tatttttccg caagatcgat tagaaccgaa caaaagtcgt ttaaggtgtt gaggctttaa 300
 acgatctttt tgattttgaa aggaggggag cactgttaaa gcgctggacc tttaacgata 360
 tcttggtttt gagaggagag aaatgttaag gcgttgatc tttaacgatc tcatggagtc 420
 gacaaaagcg gagctttggc tcctacatat c 451

<210> 30298
 <211> 503
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30298

aggcatacga tcangctntc gnananacac agaatnaaat ccncggnncc cgagactccg 60
 nnggagngga cctgcacgca tgcttgttta tttcttncna accaancggn gcgggataca 120
 gtgtggagtg tatagacttc acagcataaa naataatcag tctatgttct ctcatacatt 180
 accgcatatg gagatgagct atatctcgtt cacataagac tggacaatac cgctgtccat 240
 agatatgtat tatgattaca aactcgctta ctgaaacctc tctcgcgaaa tgagtctcta 300
 cattgattaa ccatctacat aatggaaata gaatggagag atgtctagaa atcagtgcac 360

catgccgcct atacactcgg agatcttatt cgatggctta ccctactata cctcgcacag 420
acagagtatg gtccttatct ctgcgggacc acttcatcaa aatgtcaagg agctccagat 480
actcatcata cattcactca cgc 503

<210> 30299
<211> 398
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30299

aggtttctgc ttgannccan anataacccc gagccnncga aggtagagag ctctagtgtt 60
tnccaaaacn gagagacgag ngtgaaacaa acacaccccc caaccaggcn gcaccacaaa 120
ggaagagaga aacgccagcg gagaccgaac gcgagatgga aggatcagag gacgttcacc 180
aagcgggctg gatttgaatc attcctgagg aagaagatga agctcttacg aactgtgtgg 240
ggtgatacta catatcagta tgacaaatca gatcggcata ggatacgcca catggaggaa 300
agctcatcca ctggaagaaa ttcgtcaaag aagcaagctg gatgtagctg tcccacaatg 360
aagctggagg gcgtggcttg gggatcaa at caagcaag 398

<210> 30300
<211> 415
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30300

ctatcttact ttttcttgta tcgtgactct tcgttgccat catagagagc ggngcagaga 60
gaagaaacac tctctggcct ctcatcttca agcttcgatg gagatgagcg ttgcaaggct 120
aaagaaggac gagatccana ggctgaagaa agagatcaat tagctccgac gtccggcgac 180
agagctgcat gactcagaga caagcgcgac gctgaagaac ctctcgaag agggagaaag 240
aatggtgaca ttcctagaga cgagcgcgcc agcaccacca tcaccatcgc tgatgttatt 300
caaaccctaa cccttctcac cctcaccctc aaccggtnt gctcgtcttt tctgtggga 360
ccgccttcaa cgacgtcgtc gaggagctca agaantcac caccgcgac gctca 415

<210> 30301
 <211> 408
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30301

ctaataaatc tatgtatgat ntanaacaag cctcacgtca gagttaçctt tagtttcatt 60
 ggaatatctt cttcttttgg ttttgaggaa acccacatgg atcaatgcat attaccacaa 120
 ggtcagtggg agtaaaatat gttttcttgt tttatatgta gatgatattt tacttgcaac 180
 caatgatcaa tgtttgctac atgaggtgaa acaatttctc ttttagaatt ttgacatgaa 240
 agaatttggg tgatgcatct tatgtcatca gcatttaaga ttcatagaga tagacctcga 300
 aggatttttag gtctatcata ggaaacctat attaccaaatt tttatagtga ttttggatga 360
 taattgtcac caagtgttgc tcccatcgag aagggtgata gatttaatt 408

<210> 30302
 <211> 372
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30302

ttgcacctga tggctctgtg ttttcaccaa ctttaattcta gggtgacaag ttacatcctt 60
 gttgattcga tgggttgcgac tcagttcaga ttgtcacaat tggcttacga gatttgaaac 120
 acagggttaga atatctcaaa ttcataanaa tggggttatg gggttttcgag attatgacta 180
 gaacatgaaa atagattaga aagaaaagggt tccatttttc ctctttctaa gttgaaatnt 240
 agtgctgcta cctttaccct tttcccaatt acccttgaat taccattttc aaccggattt 300
 caaactcgtt ctgtttatct tctctagtta cataaccatt gctgacaatt gtgtagtgaa 360
 gtattaattt tg 372

<210> 30303
 <211> 445
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30303

gtgagttttt tctttttgat tgcacacnat natgtttcgc gcaacanncc nnnaccggag 60
 gatgcaacaa ggtgctacat tctnaacncc cggaccnngg ggnncggaaa gaggggaaac 120
 accccggggag aaaacgnncc cccccaccn ngaaaggacg acaggggagc acgccaggca 180
 gctgccggac cgcgagaagg gggcacnacc acgccacccc cccgaaggga agcggacagc 240
 gaacgagAAC ccaccaaagc agaccaagcg gccgagacgc ancaaaaaca accaaaacac 300
 aaaaaccgca aacgcanagg agactgggca gccangcacg acaggcacgc aagcaagcg 360
 cgaagagggg cacagggccg ccccgaccaa cccaagcga accaccaag accggaagaa 420
 agcccaagaa cccaccgcg aacgc 445

<210> 30304
 <211> 464
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30304

aggatttctt gcctangatn ctnannantc aagacccgcg agcgntaaga ggccggatgc 60
 tgcaagcang attgtgatat caaaaganac aaccggcgcc gcggttgat cttataagag 120
 gcaagcatct atccttcaaa cctgtgctca ccataatcga taacctacaa ctctgcgac 180
 aaacttgatg aatgcttga tcatccacct ttctaaaaaa tgcattgctc aacctgtc 240
 atttcccaag aaaagtgtta tgttcaaaaa cccgtgcata taatcgcttc atcctctact 300
 gcctatgcga aagcttaaaa gaactaacca cctgaatcct ttgtggctct ctcacccttg 360
 ccgaagaaga gcgacaaccg cctgatgctt tgtgactctc tctacaaag atcgaagact 420
 actgctgaga tcttagaaca tctacccta aacaagacca aggg 464

<210> 30305
 <211> 566
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30305

ngggatctta cccggaattc gacgnacnna tncatanca ncnaaanaat tttctncncc 60
 gngngagtag ccctcgaggt ggagncngcg ctggatagca tngcatgcaa ttggatgant 120

tttnnnnaag ganagaacct cccggctgtg tattgtgtaa ccacacaagt ggataccctg 180
 nnagatattg tccccggggt gtcaaggaaa acccttgngg acgatcaagt tgtggcgcta 240
 tttgccnat acccagcctg accaatccca accaaccg gcataatcag tcatttgaga 300
 acctgtaatg tacctaagca ggcgatgctc tggcagtaaa cagatggaaa ggaaaacaaa 360
 aaccacaaaa ctatgggagg cttgttggtg ggctggccca actgtgaatt ctctggaatt 420
 atagtggatg gtagccctct ggtaatcnat tacctaaggg ctgggtaatc gattacaaag 480
 cctaaaaatg aaaacaggag gctatgattg tctctggaat ccaataccac ggggtgatct 540
 attaccaggc ttggaaacaa gtcacn 566

<210> 30306
 <211> 490
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30306

cggaattttt agttcttagt atcttgcacc nctnatnaca cagcagcctc tcgaagcgag 60
 agcnnaggag agaaccaca taatggtcca atcccnccca naccaccggg agaaattgaa 120
 attccaattt ttaaaccctg tatccgatta cacaattgtg gtaatcgatt accagcagtt 180
 agtaaacggt ttattccaaa tttaaaaagc tgaattcgat tacacaatgg ctgtaatcga 240
 ttaccagacg ggatttcaga aaaatagttg caagagtcgc aactttataa atgctttaca 300
 tctgaccacc atgggctatt tatatgtgac ttaacctgaa attgctcaga gattttcagc 360
 caacagagtg ttatcctctc aaaagcaatt tcatttatcc tcttaagata tcttagcaat 420
 tcaatgcatt cttaaggatt aattgagtgc tcatctgtaa atccatctcc tcaagaagat 480
 ttgttctttg 490

<210> 30307
 <211> 414
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30307

agcncatagc tctctcttca cggaatgccca ttatggagtg atcaatggat attcttcttt 60

caagtgcattg atacaaagat atggagttcc aagtgggaaa tataagttgg ttcccatagg 120
aactagagag gttgctaacc aaataggacc caacacaact attgtaccta aatctacatt 180
ctatacattg ttgtaggatga acctatcaac agataaacia ctatgggtatc ttgataacgg 240
ttgctcaagg catatgatag gagacaagtc aaagtttatg tctctaaacg ctaaagaatg 300
aggatgtgta atctatgggtg ataacaacia acggagaatt cttgggtgtag gtaatattgg 360
taattcccta actatctcca tagagaatgt cttatacgtg gaaaggctga aaca 414

<210> 30308
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30308

actttattat gctgctcctn ngctnntaat ccactgctta atatccaaaa taaacttgca 60
acctttgact tagacttgag aaagaaatcc aacacattct ggtgaaatca tctacaaaga 120
taatgtaata tttactctct ctaagtgaag tggtcctttg aggtccgcca aattttgtgtg 180
aatgaactgc agctttctcta ttgctctcca agttgattgt ttgaagagta atcttggttg 240
cttgccatat taacatgttt cacagtttgg taattcagaa tctaaatgag gtaatccatg 300
aaccaactcc tttcattgca tgtccaacat aactgcatga tgataatggc ctaatctttt 360
gcgccagact tctataatat ttatagtaac tgg 393

<210> 30309
<211> 276
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30309

acccatggcg gagcgaggca gagcgtcgat tccttcttgn cagngnnaga agaaaagatg 60
gagtagcaaa ctaaccctag cgcagagaag agatattctt cttcgcggta atatgtgcag 120
tggcgagctg gccgaagata cctgacctgc gttcgaaaat caacccccctt ggaaatccca 180
aactagtgt ggtacctgtg ctcgacgatt ggggtgtcaa ggggaagtac cttcggcgag 240
cagacttcga cgcattcattc gtgaccttcg caaacg 276

<210> 30310
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30310

agcttangtg atcataattg cctcaatcat ttccaaagtg catgtgaatt anggagcatc 60
 aacaagaatc aagccaaggc tattgtgcaa gcaatcaatg nggaaaaaca caccacatga 120
 ttatgatgat ggatggctca aattctcaca atggtaaact catcactttc aaattgagct 180
 ttcaaaacta tcatgacatg tagaggagaa tcaaggattt caagtcacaa aatgtcaaga 240
 acttttatta tcaaaacaat taccattttg ttgaacatat cctataattc atagaaaaac 300
 atgcaaagtc gtacatgcac acaaaattga ccataatat taaactagaa atccgacgaa 360
 actaacaaca ttaacaaatt aacacaacta acaaattaac aaaccaacaa tactagcaaa 420
 ccaaagaca 429

<210> 30311
 <211> 449
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30311

ttgagacaaa tgtggaatgn tgagcactnc aagatagatt ccggctatct ccatggggca 60
 agagggtaat gaaagaactt ccaatgacta ctcataacat ataatgatct gcctcgctgc 120
 tacgatatca ctactctaaa atgagaaatt tcaattttta gtgaaagttg tattaatttg 180
 attatgaaaa tgggtgagaat atttttgcga tatacattca tcaagtaatg catagattca 240
 cacacgcaca cgcacacacg cacgcacaca cacacacaca gacacacaca cacacgcaca 300
 cagacataca tatattaaac cactatacat cattcacatg acaagatata attcagtgtt 360
 cacatgtatc taaacttgta attgcatgcc cactcaacat cagtgaccaa ctaggaagaa 420
 ttgaaacaga catactctct aaagaactn 449

<210> 30312
 <211> 420
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30312

agcttactat attgcaattg gatgcttccc gccaccagtg atcaagcctc tttgatagcc 60
ttaaacaatct ttctttcatc aatgtgcaca ctataatggt ctctgaaatg ttcattgtggc 120
tccacatgat ntangtttgg atgaaaccta agcttatcag ccaccctttt ttccatccat 180
ttcattgtag cttgtttatt nttgaagacc cttccatata tgtgctcctc caaaaaagtg 240
ttgatttgaa agcttcttgt aacttcgaac catgaacaat aaatttccca tgaacatcca 300
acttgtttac aacgcgctct agcttgaatg ttgtcaactt taccatttcc agatctcttg 360
catggaaaat agttaagtct ctaacaactt caataaacat tntgatgcta tcaaactcca 420

<210> 30313

<211> 455

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30313

gccgatttag ttnttgctcg cgagaggatc gaagtttggt ttaattgttg anaatnngat 60
natectactg tgatgattgg gattcctang gcanatggag agagtaagaa tgagggagga 120
acccatgcta tgactgccat tctacatgg ccaaatttcc caccagctca acaatgtcaa 180
cactcagtca atatcagctc ttctcattac ccaccatcct atcaaccaag aacacccaat 240
catccacaaa ggccaccctt aaaacaccaa ccagagaaag aattttccag caaagaagcc 300
tgtaagattc accccaattt tgggtgctga tgctaactta ctcccatatc tactcaataa 360
tgcaatggta gccataatcc cagccaaggt tcttcaacct ccatttttctg aggatacaac 420
tcgaatgcaa catgaactca tcatggagga gtctc 455

<210> 30314

<211> 326

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30314

gggatgcctg tntcncnnnn atagaagtcc ccgacgngag gacggaggag cacttggtct 60

tgaaaccaag gccactgttc ggaatccaca ctgacttcaa agaggaggcc ctctcataca 120
 tgattcaacc tccccaaaca atattgctag gtcgaacccc gttggactca actcccacga 180
 tcctacatat aagaggacac aatggagtct agtgggcatg atcacacaac gtgctgaagc 240
 acgaagatgg actgcaccat tggaccttac cctcatgaac ttaaccaggc catctaattg 300
 atagcccata tgactgaaga gaactg 326

<210> 30315
 <211> 387
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30315

gacctaagta aactaactcg cttagcacga catgctggct tancgagtcc atacaaactc 60
 agaaattaaa aactagaatt ttaaactctc gttaagccga agtacagtgg cttagcaagt 120
 tcatacataa aagcataaat tcaaacataa atgatgaaca cgcttatcgg gacagggctg 180
 gcttancaag ttcattcagat aacccagaaa ttcattccaaa attgatgaat tagcttagcg 240
 agtacatcga aatttcctaaa aaattggggc ttgaagccc ctactttcca gtcactttca 300
 ggctaagaa ctctaataca aacacatcaa atgaacctac attacctaag aaactagatc 360
 cctaacaaca tataatcaaa caactag 387

<210> 30316
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30316

cagggatgcg cctaattgcat gaacatattg atattaatgt caagcatttc gttcttcgag 60
 atgacgatcc tgagaacttg aataaattac agcccagaaa tcaaccaaatt tagtgcgatc 120
 catcttgctc tttaattaat catccactgt ggcaatatga tccacaatta gtggggtaaa 180
 gtttatacac aagtcagatc aaaataagag aattntaagt ttatgcaaaa cattggatat 240
 tattcctcan aatatattan aatgaatgac atatattgtg cattctcgtt gtgaagaata 300
 acatttcctc cactgacacc tcatgtatag gttgcagcca gcatgcatgc aatgagatat 360

atatggatga aagcaacaag tacgtgtctg agtngaactt gaacta

406

<210> 30317
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30317

nggggtgtatt ctttnatctt gcannncgcg aancaacaaa cncaggcttg gaagagaagc 60
gttgaggcca ttgattgata ttcattgtcc gcancggccg ctgagtgtc ttgcgtctgg 120
acgagaatga cccactgatt cttcctttgg tggagacatt gttcaggcgc aaatgattca 180
gtggatggct cgactcagag gaagaagatt cggtagtgt cgtttcatta gggagctact 240
ttgaactttt ctaaagacaa atggaagaaa tcgcccttgg gtattatatt ggggacgtcc 300
atcttgtggg tcgtaaataa taagtactaa ctgaaataag aagaggagag ggacttggtg 360
ctcgaaagaa ttggaaagtg gggagatagt acatgtgtct ctgtggaggt cttccatctt 420
ctggggtgtt attaccctt ggggtggaac acctggaagc ctgtt 465

<210> 30318
<211> 324
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30318

ttttttgttt attggcntac tcaaacagg accacgtttc atattttctt cctagcatng 60
acataactgt cagttactgt cgaggcttct ggagcatcta taacttggtc actatattct 120
gtgcgacatt tgcgtggata agctgcatca aatctctctt gtctctctgg attccttcag 180
cacgaaagta gtttatggtt gtcaattgct tggcaactgt acttcgtatc tctactttga 240
caatcttctt ccgtatcgca tgcagagaag ccccttctc aagctcttgt aacaatactt 300
ttcttgcttc ctcaaatacc atct 324

<210> 30319
<211> 618
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 30319

cgaagcgatg acnactangn ctctacgaan nncgcgcaca cnatatgaat aatcaatgct 60
 tgnnaggatt atgggagtag gccgatcaca tgtgtgtact attgtgtgtt ggctcgcgcg 120
 aatggtgcac acgcagagtt tatccgacat attttatant gcggcacata gaacaccaca 180
 natcgcccggt' gtgggccgca cactaccaag ctggagcgtc agcgtaacctt cccatcgtag 240
 gcccgaatat ctctcngtct tcgtcatcag acacgcagag ggccgctcat gcanatctcg 300
 tcgctcagag cgtatccgcg agacaatcgc gaaggtagat ctcaaactat tgcaagacag 360
 ccacaacacg tatcacgagc gcaagaaaaa catgggcgaa agagcagaag aactcatgcc 420
 ctaaaactac caacgcaaaa gtcacgagct gggtccacg ttaaaggacc gccagtgage 480
 atttcttttc gatccaagtt cggtaaccag ctggatcgac tcagtaaattg ttactggaag 540
 tctctactac aaaagcctac attttgaccg ttgagatatg ctagcacata tccagaagtc 600
 attctgcact actctttt 618

<210> 30320
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30320

ctagccggtt ataaggcaca tttattctga aattntaatg gaatgtttgt tgatgttatg 60
 tgctntaagg tttttatttt cgtttattta taaaataaaa tctgtccatt tgtatgaata 120
 gacaattatg catagtgtga acaagaaaaa aaaaaaaaaa aagagacact tgtgcaaagt 180
 caattcaacc attgtatctt tttttctcat ctagaagttt gcatagattt ataagaaaac 240
 taaaaagaat tagtgcaaac tccaaaattg atccttcagt ttttgtcatt aaattagttt 300
 ctcaacaata gataagaaaa aaaaaacact aaaattcatt ttcttgtgtt acaaaaatgt 360
 tcagaggttt aagatgtgaa gtgtgaacac tgtaatctt 399

<210> 30321
 <211> 139
 <212> DNA
 <213> Glycine max

<400> 30321

tcgcggactt aagtcaatgg tcaaaccttc accattctac gcttattcca ccaccttggc 60
cgagacctcc cctagggcac atgtccttac catgtgtgct aagacatgtg tgctttcctt 120
ttgggcttct actgagtct 139

<210> 30322

<211> 419

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30322

agcttggtgt tctttgtgtt tgctggtggc aaatgggttt attgtattac atatgatgtg 60
ggggtgttag ctcaaacctt gtgtgtcaaa ctcatatctt tatcaatata agatatttgt 120
gtgcaaaaaa aataaaataa tgaatattaa aaatcaccgc aatattaaaa gttatttcaa 180
aatttaaagt ttaataaat attaattntt taaatgaaaa cttagaaagt attaaaataa 240
tacaattaaa aaatataaat aattaaaatg aaaattntta aactcaatat attaaattga 300
aactaaagta aaatttaagt taccaagtgt cattaagtct tttaaataat cacttaaaaa 360
tatcaattga tgagatttca aataaaaaata ttacatatt tgcttatgac anaaataaa 419

<210> 30323

<211> 479

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30323

gacacaatat aatacagcag ctgtggtgtt ntgcagtctg tttggtggca ctaaagacat 60
ttgttctccc cttaatctaa taggccttgt ctagattgtc atgttcaatc agaaaacacc 120
aggtgaaatg gagtctggaa ctctccacac ctataaacag ggctagtaaa ttttggtttt 180
ttgttttatt tacgaaagat aatggatctg atggagcaag cagatataat ggcacaacaa 240
gttttccga aggaacttat aggttttagtg tcttaataa acaagatgtt aatcggattg 300
cctgttatat tagaagtga ttgaattcca ccaagtaata gcattcatag gtaatgttgc 360
acagacaata aattaaaatg tatattntct gcctttttct tttcttggaa gtcgatgcaa 420

tgacatgctc ttcactgtgg tgagtaaaag ttcagataca tctacatagt atactaata 479

<210> 30324
<211> 479
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30324

agagcgggta gttcttgaga tcnegcaata ntgctctgta cccgcgatcc tntanaatcg 60
aacctgcagg cttccaacct ggactattcc catcccaccc eggccttatt cggccgtgga 120
gaccttgtat gtaacttaac cagccaacct cttgccgtcc accaattaaa tggaaaccag 180
aacaccaaac cagggagcct ggtgtggcct gccacctgc aaattttggt tattatgtga 240
atggtggcct ctggtaatca ataaccaagg gtgggtattc gatacaaggc ttaaaatgaa 300
gacagaggct aagatggctc tggaatcggt accacgggtg taatcgttac caggcttgaa 360
acgatgtcan gaagctatga agcctctggt atcgatacca agtgtgaatc gataccagct 420
tataaagaac tggagtgatg aacctctgaa tcatacagcc tggatcata cacaagaag 479

<210> 30325
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30325

agagatgact tgaacgcca aaaaccgctg tgagactttt gaggcgtact tactcatggt 60
cttatgggat tcccctggaa tcagcggaga tataagatat atgtgagagg aggcgccatt 120
cctttaggaa taagccctgg gagaaggac cttcccacca cagatgaagc cttggatta 180
agaaagcttg gagaaagatg cttccattg gaggaatg aaagaaggga gagaaanaag 240
agaggggagg agcctcgana cttgatggaa taaaagagg tatagaaatg gaacttttga 300
agtatgtctc acaagactct cattcatcaa agttacaaca agtggtacac atgcttctat 360
ttatagacta ggtagcttcc ttgagaagct ntctagagaa aacttncttg agaagcttct 420
ttgagaatac ttccttgaga agctagagct tagctacaca cagcctctc ataactaagc 480
tcacctcg 488

<210> 30326
 <211> 219
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30326

ccaaaaccag cttgaccaat cccgaccac cccgggctta gtcagtcagt gagaaccctg 60
 tgatgtacct aaacaaggcg agctcctggc agtcaaccga taaaagaaca aagaccacat 120
 agcaaggggg cttgtgtggt ggctggcaag ctgtgaatct tgtgtgatat atgggatatg 180
 gcctctggtta atcgattacc anaggtgggt aatctatta 219

<210> 30327
 <211> 464
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30327

agaagcattg ctgagactgc accaaaaacn cagcttatag tgcggtcttg gagacaaagg 60
 tcagnggtcg ccaatattgn agatgaggtc ccaagtcctt cggattgggc ccgaccatgc 120
 cctctgatnt ccactgggaa attggcgaag ggatgaaccc ccccggtctt accccacaag 180
 cattatgtaa cccttaccgg ttttaaaaac cctataatct ggccctagct ttagaagttt 240
 catttagtaa aggcttgtgt ctttggtttt gaattatata ccaagatctt cttcatctga 300
 tcttgtctct accattctca ttctttgcat gttacttctt ttctgaccgg cagattcatg 360
 acgagtcacc gagagactaa tacctggacc cgctatcaac tcgacaagaa cgatcaacgg 420
 agatgaagag agagatgtgg acttcttcga ctagaagggt gccg 464

<210> 30328
 <211> 431
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30328

cgggcccaga aggagatttg accttaatct tgcnnattca tagtaccngc actcaagagg 60

accgaggatg agcaaatttt tcttaangac ccaacccaaa gatttctcta taccttgta 120
aatctaggaa acctatgggt cagggcaatt tactctaatt tggggaagga accattagaa 180
tgaaaaggaa aaggtacat tccccccac aaataagtgt ttgttaaaaa agaagcaaaa 240
aaataattgt gtggtaccaa aggtgaaagc acttacgaaa tgaataggag aagctattgt 300
acaaaacaga aagacattgg attatctaga cttggctctc ttaaactaac gttgaatcta 360
aaaaccagga tttttgacca cacctcctac acctgaaaat cttctatcta ttatatTTTT 420
acttatgact a 431

<210> 30329
<211> 341
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30329

agcttttata ggatttattg aanatctcga ctntatgtgg gcatggatgt ccgtagacg 60
tattatatta caattagat agaagtcttg ttattaacct catctattat tttattagaa 120
tcatcttttt gtgtattatc ttattagaat ctctaacttt tttaaaaaaa caaagacatt 180
ctaagatggg tctttgaaaa accatcttag aaagtataca ttctaaaata attnttgaaa 240
aaattatctt agaattctta atatgtttta tttaaacaaa aacgttctag ccattctaga 300
aaatatacct tttaggaagg gtctttgaaa aattgtctta c 341

<210> 30330
<211> 489
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30330

gagtagtta gtntctgaca caanatactc acagcggagg aaagaaggcg cacggataga 60
ctgcattgta ctctggtag agactgatta atggggcggtg gagagaacac ataagacacc 120
ttcgtatgtg cctacctata aatctctcag cgtagcccaa gagataaaga aatagaaaaa 180
ccatgtttga aatatgtctc ctaacgtctc tttactgtcc ttacatgtac ctactatcca 240
tgaggatacc ggatgccgat aaatagtctg actaattcag aaaatgggac agtccgccga 300

ccagttagt gatcgagttt cgaagaacat cattcgatgc tgagagtact gtatgaccgt 360
 ctccgaatga acatgatgcg tgaaggggtga atggcattat gagacgacct acaggcaagt 420
 tgcacgagg agacaccttg gcatagtggg gtgatatcaa gggataatca accgattgac 480
 caagaagcg 489

<210> 30331
 <211> 397
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30331

agcttcattg atagaaattg gtctaaattc atcaagctcc tatgagccat caacttttgg 60
 aacaagtgtg atgaaggagg ggttgacccc cttgtgatta caccatgttc ataaaagtca 120
 gcanaaacct tcaacacgtc atccttttagt gttgggtcaaa actttgtaaa gaacttaaata 180
 ntaaagttat atggactcgg acttttggtta ctatcacagt tccttactac ttctctaata 240
 tcaactctct gaaacttttc aacaagcata tcatttttca caatagggtt atgtttgaac 300
 gactctntaa gccccctaa ccttgggtcta atccccctc atattgaaat ctctctcaa 360
 agaaacactt canattctct ntaaccagca ttggatc 397

<210> 30332
 <211> 448
 <212> DNA
 <213> Glycine max
 <400> 30332

tatgtagtta aaaagggttag tcttctagtc attcaaaatg tctttttggc tcaacagacc 60
 aacctatgta aataaatacc aataattcca taataataat attattgtgt gattaactta 120
 ttactataat actttaatat atacttggtta gcctatttaa aggattatat tcaactagctt 180
 gcatacaagt gtagactcat tagcctatgt agaagtatgt aacttattca aattcaatag 240
 acctttacca catagtaagc atttaaataa acttccaagc ttaaccaaac ttttaaaatg 300
 tcaagccatg ccttaaaaaa gcccatatcc aggaacaag gcagagctca gacctttgat 360
 ttgtaaagta agacacgtc aagccttaaa tcctaactta actcaatccg tttccacctt 420
 gacctgatca ttaccacgt ctaactaa 448

<210> 30333
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30333

agcttttattg taatgtttacg ctccaacatc attttgagac gcacagtctg cacaacattc 60
 atgaaaacaa cagaggtcaa cagtntaaat atccaaggct aacggagaat gtgtatgaag 120
 aataatatat ggcattgctta cgtgtaatcc ttttttttca aaatgaagat gaaaaaatct 180
 aatgtttgtca aactactatg tagcctctat ganaagatga cctcttctca gaagaaggct 240
 tcaatcaacc agagttaatc aagagaaacct aagtcccaa tatagatata caatgtatcc 300
 atgacaaact aaaatatata tgtatacata tattgatata tacatattga aacaaaccca 360
 ctageccaaa gctgcacata tatatatata tatatatata gcaataacct taagag 416

<210> 30334
 <211> 467
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30334

tataaaactc aagggatgat tttgcaagat ctacccaac agatcatctt tttcaggcct 60
 agtnngcaga gtttcaattt ccaaccaaca catgatggta tntaacattc taattatatg 120
 tggaatagga tttgtggacg atagggcgct tgtttatatt ctccaaaact tcattagggga 180
 acatgttcta atttggatga tgcccatggt gatnttccag ttaggtgata aataaaagga 240
 actgtccatg gattactgga tgaacaaaaa taagtatatt aaagtntgt atcctgctcg 300
 atgtctgttt tagtcaacat ttgagactta cttaatgttg acagtagata aaacattttg 360
 aactgataat tgggtggtgtt ctttaagctg tatgcttgca ttacatttnt cttctttata 420
 taataattag gtcatagcta tttctacgtc tcaagtgtnt attcatg 467

<210> 30335
 <211> 501
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30335

agacttctat tcattcgatc acacnenaat naagctcgng cgcgggatcc tctgagncga 60
cctgctgcat gcaagcttgg ttcnatttnc tcaacatang caacaacgag cgatgggtcat 120
tcatagacca ctcacccaaa tatctgagtg cgctgggtac atcaaagat tgtgctcacc 180
tctttcggcc gaatcgatta tatatatata tctcataaca aatcctacct tcgtataacc 240
acctgctact agagttgcag gtttcctcaa cattgctaga ggacatatgc cagccttctg 300
atagatctga tatacaacgt atctgcttcg ctaattgttt atgacctggt attgaaccag 360
acatatgtgg cgccatctac atatggctat atgcagtgtt gaatacactc acctgccatc 420
aatgagatag ggtccttcag tgttaccgct tcatagcttt caatcacatt tacggccatt 480
tttgagttat gagtgcgtcc g 501

<210> 30336
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30336

gagtatatac tgagtactcc ccnctaacac naagctggag atatggcggc accataggca 60
atatgggcca cttttgctcc attctagcca tcangacata aggaggcccc tcttgaccat 120
ggaactccat atactgacaa tgtgaagaga gctataggca gcaatgggtc acacatgcgt 180
gatctataat gtccgaagca tggacagatc cggaacaaag actcatcatt aattttttga 240
ttaactctcg agctggaacc atagtgttga aaactgttaa tggctcgaac tttgttaata 300
caggggaaaa tcttttccaa tcgcctgatg ccgctggaga cgaagctgtt aaacaaatgt 360
tattctagta gaaaccaaca atgagaacta ctaatgttta tccgggaaat tggtggagga 420
aacaaggaac atatctattg actccgtgag cactcattga tctaattatg ctccagatat 480
tggaaccg 488

<210> 30337
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 30337

 attacctttt acctcgggtg ctgaatgaag tcccgtatat attcaaacgc ctggaattga 60
 attcccaaac tttgaaccaa attccaagac cattaccttt ttcctcgatg gcagattgga 120
 gtccggaata tatcgagacg ctcgaaattg attattgaac ctcaagcana ttcaaataaa 180
 cataacttttt actcggatgt ctgattcagt cccgtaatat atcgagacgc ttcgactaga 240
 atgccgaaac tctgaganat tcaaacgaca ataactntnt agtcagatgt ctgattcaat 300
 ccccgatatat atcgagacgc tcggactnga aaagccgagc tctgagcaaa tcaaacgaac 360
 aaaattntta ctcgatgtc agattgaggt ccgatatatn 399

<210> 30338
 <211> 437
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30338

 gaaactcagc ttcacattca attcaagcgt gtcgatatat tacgggactc tatcagacat 60
 ccgagtaaaa ggggttattgt cgtttgaatt tggtcagacc ttcggtattc catttcgagc 120
 gtctcgatat attacggaac tcagtcagac atccgagnta aaagggttatt gtcgtctgaa 180
 tttgtctcaga gcttcaacat tcaatttcga gcgtccggat atattacggg actcaatcag 240
 acatccaagt aaaaatttat agtcgtttga atttgtctcag agcttcggta ttccatttcg 300
 agcatctcga tatattacag gactcaatca gacatccgag taaaaaatta ttgtcgttcg 360
 aatttgtcga gagcctcaac attcaatttc gagcgtttcg atatattacg ggactcaatc 420
 gaacatacga gtaaaaaa 437

<210> 30339
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30339

 tatcttatat ggcattccaa tctccttaat aagagatggg tganaaataa aatgatttgt 60

gaaaatcagt ggatctgtat ggatgatacg aaattgcacc atttcaattc attcttgggt 120
 cataagactc agccatccaa gattcctctt atgatatgaa tcaatagcgt tgccaaattg 180
 aataaaaaac atgactcata agattctaata aactaggatc ggctgctgcc gctatccaag 240
 atcagtaact atgcgttaaa tactccattg gattgcagca cccaattaag atatcaagcg 300
 agcattgagt agacagttgt caataattta agaacaagct ttagaacaaa ttatctagtc 360
 caacaaggga agattgaatt taacgagaag aagaagtgag aataccttga 410

<210> 30340
 <211> 520
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30340

gggcagtagt gcttngngang cttegcannn ncgcgacacn ataganntac tcagctntac 60
 acaatgantt tattataaca cataatcatg ctattattat aaattgtgga aggattgaga 120
 gtatagtgga ttagttcgat gaatttcattg aaatttatat tggttcaata tatttacatt 180
 taacctctat tatataatac tgaatatattt aagcagtgac aaaatgctta ctatcaaaat 240
 aaatatattt atgcctgatg atgcttacat gtgagaataa gcatgtgaat aaaagaaata 300
 cgttataggg attcataaat tcctaaaatg tatattttta gattttcact cgtttgctat 360
 tctgcttaaa taacttgaca gttggacacc tggacacca ccgaaatttc aatgctcttc 420
 ataagtcgac tattaaatgt gtcgcgcgtc actaatttat ttagattcta attctatcga 480
 aagaaccact gtgttttaata cattctataa tagatatagg 520

<210> 30341
 <211> 360
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30341

ttttcttctt ttaaagaata aaaatgaang aaggtagaag agattataaa aaaaagatt 60
 accttcagag naagaggcct ataataatat ttattnttta ttttcatatt tactctaatt 120
 aattcaagat aggtcatggt atgcattcca ttgtccctct catntcacat tagaagggtt 180

attcttatca tgcacaacat aatcatttat tttaggtaat atctaattatt tgataatcaa 240
tatttttttta ctatataacc aaactcaciaa ttcanaatat ttgataagaa acatagaaat 300
caaactaata tataaataaa aatacccaaa ataaactatt agaaaaatcc ctattacaca 360

<210> 30342
<211> 324
<212> DNA
<213> Glycine max

<400> 30342
acctataaaa ctcagcttct cctccggtat cggttcaagt cgattcatcg ctcttatatc 60
ctcgccctcga tcacactcca ctccgatcat caacgccgctc ttcattttcc tttacttggt 120
cttcgggtttt tgcgtcggag ggagaatcgg aaccgaaaac gagaagggaagg aggtctcttc 180
ttctatggaa gggagtgaga gtgggagagt ggccggcgggg actacagaga agggagaatc 240
ggcattggaa ttgaaggcca gaagcgggaat atgaagtgtg ggaggggtga atggaggagg 300
gggtgagggg acgtgcctct ctcc 324

<210> 30343
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30343

naagcttttag tagtagaatc atgggaccaa ctcattntat ttcanaaagg aagtcatatc 60
tagtcaaggt ctgagagacc atacaagtnt cctaacgatn tctaattatg tgggccatta 120
agtctatcat atgctgacaa tagccgagaa gcccatgaat ctcttcnggg cggagtangt 180
gtctgccatc gccttggcct tggctaacaa tcggggaagt tcttgactcc cgttcaaggt 240
aagagcaaac cgatccatcc acatgggtgc ctcttggtgt aaagagtcga tcacccttcc 300
tctagcctct ntttccgcgt atacttgggc atactcatcc gcgattctat gctcgtgggc 360
cgtggctaga cctaactctt cttggtactt ggcaaagagt cgatcaccct tcctctttca 420
gaaccatgc 429

<210> 30344
<211> 467

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30344

gagcagtgtct agatcctggc catcaaactc agctnataag cgcgggttcg ggagaccaa 60
 ggcaagccgt cgcattttgc aagattatat cccgagacct tgggattggg tccaaccatg 120
 ccttcttgat ttccagctga gaatatgggc gagtggagga acgccccggc atttacccca 180
 caagccta at tgtaaccttt accgggttaa aaactctata agtgggcctt aggttttagag 240
 gtttttcttt tggttaaggct tggggctttt gtttttgatt tattatacag agatcttctt 300
 catctgtccc tggctttacc attctattca tttgatgtta ctcttttctg aacggcaatc 360
 gatgacagtc cccgaagact aatcctggac ccgctatcaa cttcacaaaa atgaataaac 420
 ggaaatgaag gaatgggatg gggactcccc agactagaag atgggtcn 467

<210> 30345
 <211> 408
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30345

agcttgtctct atctttttct gaattcatgc ataagatact gacagttntt tcctanacat 60
 ttctaagaac gaaatagaaa ctgaaagata tggatcgat aatcaattta tcttaatgat 120
 aaagtctaag agaataataa gaatctagtc tggatcttct ttttacctat ttaatatntc 180
 taaatattat gagttctcca tttagttttc tattcttcag agacatgtaa acacaataaa 240
 taaaagactt gccaaagcca aaatcatatt tcaactataa ataaacaaga gttgcgaaca 300
 cgcgtgtaga taaattgacc actaattctg gttcgaagac cactaaataa attgacgagt 360
 ccactaatta aggagtgtct aatcgtctag ggagtataat gtatagta 408

<210> 30346
 <211> 335
 <212> DNA
 <213> Glycine max
 <400> 30346

atagcaatat gcgaaatggt tttcctaact cctacttact agtatgagat gctgttttac 60

tcaagatgct gacaacaacg agttagtggg ttatttcata ataccctaata ctatctcttc 120
tcttcatcgc aagcgacaac atagcgtaaa atagtctatc ctatgagagc atcacaaggt 180
ggactcaaac tgcaaaggaa gcgagcatgg tcttagcata cacggcaaca acatgtgacc 240
ttgggttggt cttggcaatg gtattggctt cacggagaat ggtgccacca actgcgggcaa 300
tcgagactaa tgaacataag cagacaacta tcaac 335

<210> 30347
<211> 365
<212> DNA
<213> Glycine max

<400> 30347

gttgaacatg attggattga ggatttgatt gacaaaatgg attaggggaa tgtgatttca 60
aatctgcact tatgcagaat ttgctgggtca aaataggtgc cagcaggatt ttaacttttg 120
tgcaaaaaat gcttgtgtgt ggttggctgt ggaaagagta ttacacaatg agttctggat 180
gtttgctagt agatccccac ggtcacaatg taagcttatg cactatagac ttccagtaaa 240
atthttggagt cgatccaacg gttaacgaat tggatccaaa gaattgtact gtgggtcttta 300
aatgagaaaa ctatgatttt ggtgatgtgt gagcaaagta tctgcctttg ctctgtttgt 360
tggct 365

<210> 30348
<211> 475
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30348

nggaggctgt ncatgcgtan tgctnacnan acaaacncaa gctnatggga naacatcaca 60
tgggcacnaa gaagtagggg gggaaaatta tgggagagtt ttncactaaa taccaaggga 120
gttagtgatt tacagaagaa gctacatctc acagaaacaa gaaccacaat agccttggtca 180
agcttcagtc actagccttc aattatatag aagggtccgag gaagacataa tacaaagaat 240
taaataaaag gaaagagggg gagtttgcta aatgggggca caagaaacct aagggggatt 300
ttctatgcga catgtaaagc catagctatt atctggacca tacatgcca ttctgggcac 360

acctaggggtg atgacagatc tgctggtagc ttgtgaacaa caacacagct aagggcacat 420
 acacaagcag gcacgtctca aagaatgatt agttcttgtg tgtggacctt ggggg 475

<210> 30349
 <211> 430
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30349

agcttacatg gtctcgatat agtgccatgt aggactgcgg ggtctgtggc aatgtggcat 60
 aactcgtaga aatgtgtttt gttttatttt cggacaccaa tatgnttact cggtagtaga 120
 attgcatttg naagttgaaa ttanaattac aatnaaggat angtttgatg aatggagcat 180
 ctgtatatan ttttgcgctt atagatagaa gcatttacat tgagcacatt ttgcttttgc 240
 cttttgctta tatgtattgg ctataagggtt ttgggttaact ggttntgctt taaagttgct 300
 ttgaactgaa atccaagtgc taattaagat ttgttgtagt gtagaatgca acaaggtggc 360
 ggagtggcat aagtctcaag aatctcanag aaaaataacc ttaataactg ctttgcatac 420
 atgtaacaat 430

<210> 30350
 <211> 513
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30350

nnaatggctg ggcataagagc anctncaaca ananaaacac aagcnnatcc ctttggacca 60
 acatgtgcca gttcaattac caccgcctgc ttcggtcttc cgtgggaaaa cgtcaccggc 120
 gtgttctccg acaccggctg gtacgccacc aaccagggtg ccgttgacgt cattcttcag 180
 cacagaatga aacaataccc atgcctcact cgcgaaccct ccgtcggcgg cgcggtcttt 240
 cgttcctttc tacgcggggt cgacatcgct cgctaccttt ggggatacaa catctcaatg 300
 cccgacgctg catcgctaga cctcgatgaat tggctcatga atagaccgga gtggaagatc 360
 atgaacggga gagaccatt tctcgttgcc ggtaagatca ctngngatnn tcggagactc 420
 actgaagaag aatcggtattg ggggaacaag cttttggnntt ttactgctgg gaaaacatgt 480

cgatgcttgt ggtgagtcna gtccgtgaac gcg

513

<210> 30351
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30351

cetgcttttc tttctcatc canaaggatt cggcctacc tttcttaatt gccagtcacc 60
ttcctgtgtc ttcttttggc aaccttcacc actgcttcaa ttcatagaaga aaaccttagg 120
cttctcacia atcttgcatt tcatttcaaa cccaagtcac accaccaatt ttccccaaaa 180
gataaaagtg gtttactggc atatcatcaa agtcaagtca aactgttcca tatgcttcaa 240
gatgagaaaa gcactactta taaataaaac ttacaatgta ntataacata gaataaatat 300
tgtactanaa ctataatcna tataactaatt atccccaaag canaaaacaa atgtcatcag 360
gaattcaaaa ttctgtgac tgggtcttgag tgcctatgt ctgcacatnc ctctcatctg 420
tcagatgaag cactan 436

<210> 30352
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30352

tggcaataaa tactcctaca ttaatctctt catgctttgn atgnnnnggt cgncccttgt 60
cacgggaagc cggaaggtcc atatcacctt cttaattgta cacatggggc actgcacccc 120
caaatgcaca agtaagaaga gataattttc cgggctctcg tgtccgtaaa atgcattcat 180
atcatgcac gcataagcat ctcttcataa catcatagtg gacatatact gcatttgtcc 240
gttatcatat tccagcctca cattntgcat gagtcatggc atcatcatgc atatgcgttc 300
aacanacttt ttgatctgca aaattgcata ccatttgttt tcatgtttgc tcatccttgc 360
gttntcctct acaaaacana aacaaaaaag ggggaagcgt gaaacttcat actacattct 420
tagtttcatg tgtaggcac cacgagccaa ccat 454

<210> 30353

<211> 404
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30353

agcttttagtc ttttcaactg cacaacgctc ttaatatggg aagagtatcc ttgtggaacc 60
 ttcacctgac gaagacactg acaataactt atcttttccct tcttggacaa agtatggcag 120
 gctgggggca agtaaatttt ctccccatca gaccttggat gcaactgtgc tcttataccc 180
 atatcagcta gatcttgacg ggtattcaag ccacccctcg tcttgccttg aatgttaagg 240
 agcgtcccaa tcacactgtc acaaacattt ttcttcacat gcataacatc aatacaatgt 300
 ctaacgtcaa gatcacacca gtacggaaga tcaaagaaaa tggacctctt ctcccatatg 360
 caactctgac tnttatcctt cttttgggtc ttcccaaata cagt 404

<210> 30354
 <211> 496
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30354

aagctgggcg tgcacanngc tgcgacacna tagaatactg cagcgtgtcg attcattcta 60
 tgtacccgta gcaggccaca ttgtgtttct tgcattacta tccacgacga gtggactggt 120
 tatacccagt gaggacgggc ttaagccatt ntacttaagg cgtgagtcac ttaactgaag 180
 atagaaggaa tctgccccga acggttgact gatattatcg cgtaacttcg ggtggaatca 240
 attgcgaccg ttcggtcgtc gcgaaccacg ttggaaagca taaacaggta gaaaacaagt 300
 atgtaatcga agaaacatct cgttagtaaa tagtgcgga agataatcgg acggtttctc 360
 tttgggatgt ctcatgctta atcgagttga ttggtactaa ggtgaaacta gagttagatc 420
 aactcgctag gcagctcgcc acaaaagagg cttggaagtt gcgttttgat gctcctaaga 480
 aaatggagat gtgacg 496

<210> 30355
 <211> 386
 <212> DNA
 <213> Glycine max

[illegible]

<210>	30356
<211>	380
<212>	DNA
<213>	Glycine max

atatcttaag	ctgtagctac	aaccttgatc	tccccctttg	gcgtcatcat	atagccaaag	60
aactcggaga	tgagcacagt	gataacaatg	gagtagcaag	atataagtat	cagagtatta	120
aatacaataa	gccaaactca	taatcaagaa	ataatgaaac	cagaatttaa	ataacataaa	180
atgtcaacaa	ccacaaaata	tccaagactg	aaatgtaaaa	acacgagata	aataagcaaa	240
gtacttagca	taataatgta	aatgctaaga	aactaaaagc	cgaaatacac	ggcgtataaa	300
agataaataa	tcagaatcta	atagcttaga	agactgagga	aggggtggaa	gatcgaaact	360
ctgacgaatg	tatccgacat					380

<210>	30357
<211>	361
<212>	DNA
<213>	Glycine max

12664

gcagctggct aaacacatat ccaccgctaa gcacagcttg agcgcgctta gtgcaaagga 240
gaatttggca gagcatcagc atcaaagtcg cgcgctaagc gcgggatcag tgcgctaagc 300
gcagaatgtg ccttcagcca ggctaagctc gagactggcg ctaagcccaa tttacttac 360
t 361

<210> 30358
<211> 473
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30358

actcagctag aaagcaactg gatgcgttgg tcaacttggg aacctatctt gttcttgaat 60
cagaaatctg tacctgtcgc aagggtttgt ggtttgtgct cctctgctga ccaccataca 120
gaccttngcc cttccatgca gcaacctgga gtaattgagc aacctgaagc ttatgctgca 180
natatttaca atagacctcc tcaacctcag cagcaaaatc aaccacagca gagcaattat 240
gacctttcca gcaacagata caacctgga tggaggaatc accctaacct cagatgggtcc 300
agccctcagc aacaacaaca acagcctgct ccttccttcc aaaatgctgc tggcctaagc 360
agaccataca ttctccacc aatccaaca cagcaacaac ccagaaaaca gccaacagtt 420
gaggccctc cacaaccttc cctcgaagaa cttgtgaggc aaatgactat gca 473

<210> 30359
<211> 302
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30359

agcttgnatc ggtttttgaa tacacgnat actgtttagg acgtgacata aatatacaac 60
atgatatt ataaattgat tctaaagtcg gacggaaatg caatgagcca gtttatgaaa 120
caatgtcata actcctgtgt atatggttat tgcattgtgat gattgataaa tgactccgaa 180
gtcttatact ttcgattaat atatacaggg tttgatactg gcgcttctta tttttatc 240
aatctggatg taatgatgta tagngtatac gttgacttga gatgtttgcc actaattaag 300
ca 302

<210> 30360
 <211> 380
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30360

 gagccctata ggggatggac cttttcatgt tacggagatt attattatTT atgcctataa 60
 gnnggacctc cgagaatagt atggagttag caccacttat aacattgctg atgtaattcc 120
 ttttgcaggt ggagctgata ttgatgagga ggaactaaca gatttgacgt caaatcctct 180
 tcaaagggaa ggcgatgatg cactcctccc taagaaggga ccagtctcta gaaccatgag 240
 caagaggctc gcagaacatt gggctagagc taccgaagaa agccctaagg ttcttatgaa 300
 cctcaaggta aatatctgaa cccatgggcc aagggttgcgt ccaattatct ttgtacatat 360
 tagactagga tgtcattata 380

<210> 30361
 <211> 516
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30361

 ngattgggag gatcnnagcn tcgcannnnc catnatatnc ggaaagacac acgggggatcg 60
 tntagagtgg atgtgcacgc atacttgctt ttatttacca aacngnaaca gaagcgcacc 120
 gcatgcagca taattnttat tgtacacgaa gtntcaatta gacaaagtat cgttcttatt 180
 aagaagactt gaactcattc attcctanat cctgaccnag catgatccta atgatcaaga 240
 aatgcgctcc tatectatca ttcactaaaa ctggatttcc taciaaatata accacacata 300
 caagacaggg aaggtccaag gttatattgc ctgatgcccc gagaacagtg agtgtatata 360
 taatcatacc taatgactga tccctaaaaa ggtaatacat atggatgaca agatctgaag 420
 tattactgac ttcctacat attcagtaaa taccaccatg ttggttacag agacacctta 480
 caatgcagag gattgcactt cgagaacaga cacgtt 516

<210> 30362
 <211> 293
 <212> DNA

<213> Glycine max

<400> 30362

ggcaatatac tcacacgttc gcggagacaa acaatatcgt taagttgtaa gcagtcatga 60
gtgcgatatgt ttgctacact ggccagggca gcgtgcacag gattacgttg tttgcgatga 120
accatattag taatcaatta gtaataaatt aatattttga atataataaa ttcttttaaag 180
cgagaaaagt cacattaatt attcaactaa ttgagtgaag tccttaaatg tttaaataata 240
ataatcaata atgtatgtac ctagtggatg tattatctat ctagttctta aac 293

<210> 30363

<211> 380

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30363

atcttttagta acatgcagct angaaccag aacggctgct gagttcttac agacaacgct 60
gagagaaaca aatgaaaaga taggacgata ctgctttgct tataatgttca attgaatcta 120
atgagggaat cctctttcca aaggtctccc ctctctacga gactgcaaca catcagatgc 180
ttagttcccc aatacatata aatattaatg ctttaccttc accgcttcta ctatgttagt 240
caagtgcctg ctattacaac tataaccacc ttactcattc tataacttcc tataccttat 300
tagaagcaca ttaagaacaa acaagaccct aagagctgtc atgggttcttg agtaagaaac 360
caatcactgg aacacatcta 380

<210> 30364

<211> 388

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30364

tttttttaat aaattgtctt agtggcattt acgagagctg gtggatgaca gtaaagtnaa 60
tgattcatag ataattgcat tattagaccc ataccccagc tgagtatcag ccatcagggc 120
acataacaag acaagactaa gatttgacct cggttagatc ggaatctaac ccatcaggta 180
attgtgactc ctcttataga ggaatgaacc aattaaaacc atgcataana acagataata 240

tattacatat ctacagagca agttaaattgg ctacagcagta ttcattccaca gcaccacctt 300
 ataagtgtca ctggcctcgc acatatattaa agatgttatt aanagaacaa agagaatttc 360
 attactgggc aacactagtc ctataagc 388

<210> 30365
 <211> 418
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30365

nttagttcct cccaataagt gcctgaaata tgtgtttctg atcttacaca ctacgtgctg 60
 ttngcttgat attgccatgc atatctctat gatatcatgt gatgcaatcc taccceccaa 120
 gggatttgga tagaagactc caacaagttt gcgccagaga tgcaagagaa gaccctaagg 180
 ttctcatgag ccttaattgta gattttgagc ccatgggcaa agtatgagcc cacttatctt 240
 tgtacatatt agattaagat atcattatctt ttgggccttg tatttatggc tctataatgt 300
 aagtaagggt ccctagaaat gtacgatctt tcagcccttg tattttaagg cacctatact 360
 agtttttgta ttaagggtac ttttgtaatt tcacatgcat taagtgaata tgtgatgt 418

<210> 30366
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30366

agggttttat ctttgaacac gcgatcgact cgaaccgagg acttttagtg gcctgagggg 60
 gcgccttttc cttgtattgg ataaaaaat atggggcgagg tttagtaaa aaaatgacga 120
 taacgcataa taaagattcg catcagattt tgccctctacc caaagtgtgg aatgttcgct 180
 gtctcaccag catacaatca atatatgaaa aaataggctt tatcttgagg ttgaagaaac 240
 tgatgcgtac gtggatattg cccactcgct ctttgtcatc ttctttatgc ataattggtg 300
 gccttaatag ccagaaatc aaaagcagac tccatgtaaa aaattaagac ctattatcaa 360
 ggaacctaac atttgctaaa tgaaagcttg gaactatata aaggctctaaa acacgttaat 420
 ttcatgacct tn 432

<210> 30367
 <211> 478
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30367

 cacacccacc gcaaaacgaa aacaaaccac ttaacagaac taacnnannn nacaggaag 60
 ccatatcctc ancctgcaca acaaaaaccn gaggacgaaa aagcngctc agagcgcagc 120
 angcncttat actattgagc aaatcaagca catgcagcga gacagaagaa atgaactgac 180
 aacacacccc aataatcttt ccacggagca aagaacacat atacgcagct gcaataacac 240
 cattagaaga cccaagcacg ccttaaaata aattcaagac tgaacaatgg gaaaaacaaa 300
 accacccgct gcaaaggaga aaagagaggg ccaaagacc aaagggccat accataacac 360
 aaggaaaaag cacaccactg agcaaaataa agctacaata gcaagagctc tcctacacaa 420
 aacgaaccaa aaattcagga ccaccaaaaa aagacaacca gagccaaaaa agaacaac 478

<210> 30368
 <211> 485
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30368

 agagaatggg tgaacctatt gttataccca cngncgaatt cgagcatcgg accgccggga 60
 ttctcttaga gccgaccggg agggcaggcg cagcctgttg tactatgcag cgtctctcga 120
 ngaaacacgg cggngagctt gtgcatagta ggtaacaaa aaatgcattt ggtacaaact 180
 tactagaagt tcttgacgc attctcggg agactactct atattaagct tatgctcaac 240
 taagataaac aatggtttca tagcgctctt tcattntagg gtccttacag gtcttccaaa 300
 actttgaagc tgctcttcca tttctactcc aacctatagc aaacgtaatt acgtccaatc 360
 aactaacaat attcttgccg gaaccaaagg cacgcttgga aaggataatt ggaggatcac 420
 catagtaaaa ataatatata ttcactcaat gggattatta agaacatatc ggcccttatg 480
 aaggg 485

<210> 30369

<211> 483
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30369

ccgactttga tcttgatgac tgcgcgctca tagnaacccc ccgangaata ngagtggngc 60
 cagagtatca ngacattaat ttgttttcta caagcncaan agcccagagc gcttttgatc 120
 acaaagacga tgatcttcac aatcaaagaa tgggttcaag atggaatcga taactcttca 180
 gggttcaaaag aactttggtc tcoggacccc tagactccaa cttcatgac caagttccga 240
 gattcaagat ctagaatcca gactgcagat ccacgattca tgagcatact ctttcaagat 300
 cagtttcaaa tggttttgtt aaaaactcga gagcacatga tttttcctca cacctttacc 360
 caaaaagtgt tactcctctg ttatcaatac tacactattg taatcaattc ccagtggaa 420
 aatggttgc aacagctttc acctgaattt acaaccgttc cacttggtat caaagagtgt 480
 acg 483

<210> 30370
 <211> 447
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30370

aggagatttt gctcgtatat ctanggcgat tncnagctcg gaccccgggg atcctctaaa 60
 gtggacttga ggcaggcaac cnggtgtnga acctccaang gaggcgggac ctgggagctt 120
 catggggggtc cttcatggga ttttcacatg gaaatgcacg gaagactaaa gaatataggt 180
 gagagaagcg ccattcatta aagaataagc catggaagaa tgagcttcac ccaccaagat 240
 gatgccttgg attaagaagc ttggaaaaga tgcttcaatg gaggaaaaga aagagggaga 300
 gaaagaaaga ggggggagca cgatattgaa ggaataaaag atgtatataa gtggaacttt 360
 gaagtatgtc tcacaagact ctcatcattc aaagggtacaa caaagtgtta cacattgctt 420
 ctatctatag actaaggtgc ttgcttn 447

<210> 30371
 <211> 541
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30371

gggggtcggaa gaaatcgang nnactttggn ananntctnc nnnnnntnta ngnanaagng 60
cccagcnnan canntccgcg naccaggagg ggcgagacac ggaactanat tanttcttta 120
ctcttagaag gaaatctacc tgtgcgggcc gttggtggtg gatagtaagc aaaccaagag 180
gatgcctgcg acgcgaattg gtttgcccat ttgaatntga atgttaacct taacacacaa 240
taagcatacc aagcccagta ttgggtatgg ctctcttgaa gccttaaacy gacgaaagtg 300
cataaactcc tatttggttg tatgatgatg gagaagcggc actttttgga cctgaaatgc 360
tacaacagat taacgaacaa gtgaagttga ttcgagagaa gatataagca tctcaggata 420
ggcagaagag ctatatgata gaaggggaaa ccattaattt tctggaagag acatgggttt 480
tgaaggttct ccaaaaccgg agtcagaaga gcctcaatgc tagaagttac acccaagatc 540
g 541

<210> 30372

<211> 173

<212> DNA

<213> Glycine max

<400> 30372

caacgattgg tacctcaaaa cctttacact gggcaatgag gggcattgtg cattagcctt 60
aagtgaacat acgggcaa at caaaaattct cacctgtcga tgttttttaa caacaaaaaa 120
aggagggaca aaccctatga tttaatggat tgatcaaaca ttaaaccact tca 173

<210> 30373

<211> 390

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30373

gaggataaac ttgatatcca nanataaana anaaccncgg cccantnnna gaagagggga 60
gaaggaagat aatggtcacc ccctaggcac tccggggggc aaatagcaaa aaaaacgccc 120
cctaaaaaaa tccaaccgag gcccaccgaa cgtaacgaac gaaacgcat gaatcgagaa 180

gcaccgaacg tctcgacgac cacatcacac tcaccgtctc aacaccaacg gaagacccca 240
 accaactcta aaacaccaag acccgcgggc ccattggcca tgacttgaaa ccgagtcata 300
 acctaataccc ctgtgaggca aagccacaat aagctacccg ctaccaaaaa aaaaaaccca 360
 ccacgcggaa gacaccgcac tccgaaagag 390

<210> 30374
 <211> 239
 <212> DNA
 <213> Glycine max

<400> 30374

aaccccatgg atcaatgcat atactatacc acaagggtcca tgagtaaaaa atggtctctt 60
 attttatatg tagatgatat ttacttgca gccagtgaat gggaaagggtg aaacaatttc 120
 tctctaagaa ttttgacatg aatgatatgg gtcattcttat gtcattggca ttaagattca 180
 tagagataaa cctcgagtta ttttatgggt atcacaggaa accctattta accaaattt 239

<210> 30375
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30375

ggactttttc tttttctatg aancncanna tngananatc cgncggaggg aagaggccgc 60
 ggcanncttt ttcttattaa aanaacaccc agggaggggc gggagacaaa aaaacaaacn 120
 cgggagaaaa aagccaagaa ccccccaacc gaaaagatag caaaacccaa acacgaggaa 180
 cgggacggca gaaaaaaaaa aancaacga cnaatagnac aaaacaccna aancgctaca 240
 agggtaacga aaaaagcaca catgactacc tcaccgcaac gggagaggag cgacaagcga 300
 acggccaagg aacaacgaca aaccgcaact ttctagttgt tgtatacgag tccaccacca 360
 tatatagtgg acccgactcc gaatatagcc taaacaattt tatagatagt ctatcattaa 420
 aactaatata cg 432

<210> 30376
 <211> 172
 <212> DNA
 <213> Glycine max

<400> 30376

aaaaaaaaa aacaattgct cttgcacgta tactatgggg aaaataacta ttagccatat 60
cgatcccacg aattatatgg catctcaggt taattacatg tggacgacaa aattaaatat 120
atgaagctga caataaaatt ttctccattt atggctactg tatttattga at 172

<210> 30377

<211> 284

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30377

atcttttagtg cctatcanac gttgacacgt ggcaatcaaa gctactgcct aacgggtcaac 60
atccaattgt gacgtcgggg gaccaacatt gcaatttttt ataaaataaa ggaccaatt 120
cgtgaattaa attatcgng gactaaatgc caaattggac ctaaagtang ggaccaaag 180
tgccaatttg ccttttattt atatacccn acgaaatang tactactagt tgggtgcata 240
ttaatgggtca aataatgcta agaagtttac tagcagctta tcat 284

<210> 30378

<211> 195

<212> DNA

<213> Glycine max

<400> 30378

gcaagaattg cagggttaaca tctaactgct ccaagtgaag attctctgca gctactatgc 60
tcaaaataat tctgatggta gtcattctta caactggaga gaagatctct atgaaatcaa 120
ttccttggtt tctgtgaaac cttttacca caagtctcgc cttgtatctt cttttaccgt 180
cagattcttt cttta 195

<210> 30379

<211> 231

<212> DNA

<213> Glycine max

<400> 30379

cgcaactcag cgcgcacaaa cagcacaaca cacaaaagga ctttctatag caaacgaccg 60

gacaaacagc ggcgcgacga ccagagagca ccaaagggaa ggacacgcag agcccccaag 120
aagaacacag caggctcaaa aaacaaacca aacaccccaa accaaacaga ggacaagaaa 180
agggcagaaa aaagctaagt acccgaaacc agagaaccgg caccaacaaa a 231

<210> 30380
<211> 266
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30380

ncaagctttc tcatagagat ctaggaagga taaagcgggt gaatgaacca attccgctcc 60
cgaatatgac agcctccatt ntaggagcgc tgagcaccag cagcgcttcg aggccatcaa 120
gggatggtca tttctccggg agcgacgcgt ccagctcang gacgacgagt ataccactt 180
tcaggaggag atagtctgcc ggcgttgggc atcactgggt acccccatgg ccaaattcga 240
cccacacata atcctcgaat ttatgc 266

<210> 30381
<211> 283
<212> DNA
<213> Glycine max

<400> 30381

aagaattcgc caaggactaa ccgtctgaat tcttcttggt tctctcttct cctttttcca 60
aaagaacaaa ggactaacac gctgaattat tttgtgtctc ctttctccct tgacaaagaa 120
ttcaaatga cacagtctga gaattctttt gattcttccc tttccgtaat acaaaagtgt 180
tcaaaggact aaccgcctga gaattctttt gtatcccat tcacaaagta tcacaagctt 240
aacagcctga gatctttgtc taacacatta gagggtagat tct 283

<210> 30382
<211> 310
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30382

agctnattat atcgagacga tcaaaattga acaacggaag ctctcgtgaa attaaaatgg 60

tcataagttt taactcggat gtccgattca ngagcttcac atatcgagat gcacganatt	120
gaacaatgga agctctagag aaatttcta tggtcataaat tttcacaccg aggtcctatt	180
cangcgctta atatatccag acgctcgaaa ttgaacaatg gaagctctcg agatattcaa	240
atggtcatta cttttcactc ggatgtccga ttcaagcgta tcacatatcc acacgcttgg	300
aattgattaa	310

<210>	30383
<211>	348
<212>	DNA
<213>	Glycine max

```
<223>      unsure at all n locations
<400>      30383
```

ggattntcta	ngctcgagaa	gtgaaattta	gaatgaggta	catntgaagc	aaactctcac	60
ctcacacaag	tacataacat	caatctaaac	ttgctcanac	tggatttaca	cctaaaatta	120
caccgaatnc	aaaattgact	cctcaacacc	caattttgcc	ctagaaatgg	ctcttggttc	180
actttggtca	tttgtttttc	tctctagcat	agcctaacct	ttctcataag	tcttaaattg	240
catttcaagc	taggattaac	tcattttaac	ctccatttac	tacagaatcc	agatatagcc	300
tgtcaactct	cagagcctga	ctctttttcc	actcataaca	ccacattc		348

<210>	30384
<211>	309
<212>	DNA
<213>	Glycine max

```
<223>      unsure at all n locations
<400>      30384
```

agctttcatt	gttcaattnc	gagcgnncng	angtggtatg	cgcttgaatc	tgacctccgt	60
gtgaaaagtt	atgaccattt	gaatttctcg	agagcttccg	ttgttcaatt	ttgagtgtct	120
cgatatatta	tacgcctgaa	tcggacctcc	gagtgaacaa	ttatgaccat	ttgaaatgct	180
caagagcttc	cattgttcaa	tatcgagcga	ctcgatttat	tatgcgccag	aatcggaacct	240
tttagtgaaa	agttatgacc	atttgaattt	ctcgagagct	ctcggtgttc	aattttgagc	300
ggcttgata						309

<210> 30385

<211> 495
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30385

gggcagtcta gtttcttatg ntntcannnc nnnnananta annnaccgnc cgngcannnn 60
 agaggggagcag aagaggagca cccggttgat tgatattcgc gcacaanana ncgagacgcg 120
 cgaaagagaa caanggaaac ccncgagcaa nncaaangaa canaacnnnn gcacaaggan 180
 cgccgagaca ggcaccaaag acaccgagac gcgcgaccaa gaaccacggc agcgcnag 240
 aaaaaccaag ggacccaaca gcgaacgcgg aggcgcgacg cacgcgcaa agaaancgag 300
 acgcccgaan acgaacaacg gaagcccgcg agacaaacaa gggggcagaa ccgacccac 360
 agacgggcca gacaagcgca cagaagagca agaacggccg aaaacgaaca acggaagcac 420
 gcgggaaagg caaaggacca gaccccaac cagggagccc aaccaaccng caaaacancg 480
 aacccgggaa agaag 495

<210> 30386
 <211> 173
 <212> DNA
 <213> Glycine max

<400> 30386

agctttgtat gaggtgaag aagctgttc tctgttcaa caagatccgg tggagattaa 60
 tttatctcag cctaatttgt cacaagatag tgacatagag ttgatggtaa atatttgtca 120
 caagtatagc aatataggag atagttttga tatattacaa gttttattca tgt 173

<210> 30387
 <211> 370
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30387

ctcacgtta tacctctcca tgtatcgaat agagactctg attccatang tnggaagcta 60
 gnnatcctgt ccttgatagc ttgatattc ttatttcgaa cacaattgtt tgaagaagtc 120
 gtctttgatt ctctcccatg aggttagact atggtgtggt tgagaattga gtcattcacg 180

tgcacttccc cttaatgaan acggaagaaa gtgcatctta atgtggctat ctttgattat 240
gagtattctt actaatgtgc atatttggtg gaatcgtgtn aggtgggtat tcagttcttc 300
actgctggat ccataaatt gagtgttatg gagagtatta atggccccgc tangaagaat 360
tacttgatgt 370

<210> 30388
<211> 82
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30388

agcttttatc caaaatcctg actcaccata naccttgacc cagagtgaga atgtcaatcc 60
ttaccctcgg aagcaaaaaa aa 82

<210> 30389
<211> 341
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30389

tgccaccag ctgcccag cgagctaggt tgcttcttcc ataangcacc gccttctgga 60
gaacttcta gaaggcccaa gtgggcctgg gttgctattg caccatgt ntactgaata 120
caccctttg cttttttgt tgattctttt tccgtaacgt taaagaatct tacgaattac 180
gtaacgatac ttgttttctt ttcgtattgt tatgaaacct tatggatcac gtaatcatcc 240
cttttttggc ttccgggatg ttacggaact ttacggattg cgcactaaca cttccttttg 300
actttcggca tgtctcgga cttcacgaaa tgcctaacaa t 341

<210> 30390
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30390

ggagtagacc atgtatatca ggnaatcact cgtccccgga tccctagagt caacctgcag 60
gcttgcaacc tggcttggt tggagcttct attggaggct gggatctggt gaggcttcaa 120

tgagggtccct ctaatggtga ttttccacca tggagatgca gcggaagaca naggaaaaga 180
 agtgagatga ngcgccatcc actanggaat aagccatgga agaaggagct tcaccaccaa 240
 gatgagcctt ggataagaag cttggagaga atgcttcaat ggaggaaaag aaagagggag 300
 agaaagagag agggggggagc acgaaattga agggataaaa gaggagagaa gtggactttg 360
 aatatgttca caagatctaa ttctcaaagt acataaatgt acacatgctc ttttatagac 420
 ttggacttct ttaaaactn 439

<210> 30391
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30391

aatgacgagt atgtgtaatt gtaataagct ccttagttga tattctagtc ataasnagga 60
 tgcgtgctnt aaagttttac aatgcttgaa tntgtgtgat aatcttgaat atgcatttca 120
 acttactcat ttaactttta taatattgat ccatgggttaa ggattgaaat ctttcgaaac 180
 atgttttgga aaatacttaa gtttttatcc cgcacatnat aattgattat atgatgatat 240
 aattgattat cttgatgatg atgcctttgt ttttcataat tgagaaagac tcanaattag 300
 tctattatct tgagtgaata attaattata tggaattgaa acaaatttta ctatcacaga 360
 taattaatta tatgatgata taattgatta tatg 394

<210> 30392
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30392

agggagagtg anncttcgtt actactccgc gatttcgagc tcggcccccg ggttccttat 60
 aaggcgacct tgaggcttcc aacnntttta atctttcttc ctgactactg gctctggctt 120
 tttctcctat caaatcatc gcactgtgag ttgaatatct gccctctgga tctgtaagac 180
 cactcccaag tatctgaacc cttgttgtgg attttatctt ctgcaatctc ttgagaaaag 240
 ttagccgcta aggtgatctt attgtcaaac aaggctcttc ttcaagcaaa gttccattcc 300

caccccatgc cttgataccc ttccatggat cttatgaatt gatgcttggt caatgagaac 360
 tgataaagtc taggttcttc tttggcattg acccaaatg aaatgatgcc gaattgaggc 420
 accctcattc ctttgatata ttagatttaa n 451

<210> 30393
 <211> 339
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30393

atatatatta gaggatttgt atataaatat gtaaaaatat ataaaaatca ttatccaaat 60
 taatatatat tcaaatgatt ntatattcac acatgtcttg cattatgtta atttatgcaa 120
 acatannttg aaaattatta tctttataag catattcgca tttgcatatg acttttatat 180
 atatatatat ataattttta taagaaaatt agtaataaaa aatatattac attntgtaat 240
 tattagtnt atactcctat catcataagg gggaaaagta tactactaaa ataaaacttt 300
 aaatttattg gggtttatact catatcatca agtgtacta 339

<210> 30394
 <211> 359
 <212> DNA
 <213> Glycine max

<400> 30394

acctttttta ttgcggctct ggaaaacaaa ggtcaggggt cgcgaatatg tgaaaatgag 60
 gttccaagta cttcggattt ggtccgacca tgcccctctg atttccagct gggaaattgg 120
 cgagtggagg aacgccccgg cgtttacgca acaagcataa tggtaacctt tacgggttta 180
 aaagctctat agttgggcct aggccttaga gttttctttt tggttaaagct ttgtgtcttt 240
 tggttttgaa ttataatac aaggatcttt cttcatctgt tcttggtctc taccattct 300
 cattcatttg catggttact tctttttctg aaaccgcaga ttcgatgacg agtcccccg 359

<210> 30395
 <211> 257
 <212> DNA
 <213> Glycine max

<400> 30395

gacatcaacc ggtccataga gtgtaaggag tatccacaag gcgcttctgg caacgacaag 60

aggatgttgc agaggttgga aactagtttc tttctaagtg ggggtatcat gatggacctc 120

attggaacct tgtggccttg gatcttcttc atcaatggaa gtccttgctt cttgaattta 180

atggcagcaa aatggaaaag aagaagagtt gagaggagac accacttcaa ggagaagatg 240

agtctagaag aagctca 257

<210> 30396

<211> 332

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30396

agctaatacc agtatccaga ttaagtggaa aacacgttcc aaaggtggag aatggaaatg 60

gccttctctca aaaaacctaa tttcaaccaa atatgagttt aatttttcga caatatgctt 120

attaaatcat ttaagaaagc tngaactagc tatgcttcaa atggaatcta aacacaagtt 180

cttcaaaaaca aatctaaaac atgataatag aaatcaatga agtcacaagt gaaattaaaa 240

agctaacaat agaaaaaata tattgaatca caaattctta tatagttata caaatcaaag 300

ctcattggaa aagaagaaag aaagaagaga aa 332

<210> 30397

<211> 476

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30397

nnaaacggaa cttttagttt ctacncatta aacacaagcc gaaagactan ctgctccatt 60

tacataaaca accaaaccac ttggttatat ttccgcatca tatggataga cctggacagt 120

gtatcatcag cagtaaaatc ccagcaaata atatattcaa tcctaattatt aaccggatta 180

ctgcactact tttctttcac aaaaatggaa aaattaccct ctcccatggt aaagttcgaa 240

atgagtn tac attagggcaa gtgttgccag actacttatg tatcgaggca agcaccttcc 300

caatgggtgca gtgttgctta aggcaagcat atttgtcacc taatctacan ggcaacacng 360

aagatgtaaa ggcatactg cttctataaa ccatagtagg tccttgtaa tgcctcaca 420
gacctgagat agttacacct tacggcataa caaggctatc aatctttaca gacctn 476

<210> 30398
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30398

aaaaacacgg ggcgttgtaa tnctacgnaa tananaaagg ccncggggag acccaaaggc 60
gaccagcagg aagcgcaaca ntttttttta ccaaaggnc aaaccacag gggcgggggg 120
agangcangc cacccaacgg aagacacaca cgaagaaca gggccatgaa caccgggaag 180
aacatccagt gcatcaagat ggccaacgtg acgaaacaaa gacgcaagga aagcccccg 240
ggaacaagcc aaggaacaag ccggaacgcc ggaactgcat gaccaggata tagacggaca 300
aaggaacata cctaggagaa aagagacca ccacgcaagt cgcaccattg gaaagaagtc 360
caaacaaggc ggcgaaaatg gaccaacaaa caggaaacca actgggcccg ggaaggggag 420
caagaaaaca catccaagg 439

<210> 30399
<211> 405
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30399

aataggtttt agtttctanc tnataananc ngccacgcga cgggagaaaa ccccaccct 60
tgtctttacc acacnaacag gggggggtgc acggccaccc agaggggaaa ccccacgag 120
caggggnaca ccacacaaag caacgctcaa ccttaaggaa gactgctaca aacgtcggtc 180
actcattct ccaccaggct tcctatggct ccttcaatgt agtccaccac atccatcctc 240
cataacaaaa gaggcagctc tcttgccaaa acatgcgaaa ccgagggaaa cacagcctca 300
tcttttatac ccgtcaatgc gacaatgctt ccattccat cacgatccag tattctacta 360
actgcttacg gaacttgcca aaacaccgca tatgcaatag ccttn 405

<210> 30400

<211> 350
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30400

ccccggggta ccttagagac ccaggcagg agccacgttg gtgancatcc tagacgcaga 60
 tcgggagcta tgcaatgaca ctgaggtgtc tcgctaacgc caataccttc tgatagaaaa 120
 gtatgaatag gcttagcacc ttgctcgcaa acctattctg agaaaaaat ttttcggtct 180
 cgactctcgc gggatatcgc actgacccat gactacagat tataagccgc tcatgccttg 240
 tcgcgctcac gccagtctat tctagtggat actgctttct tttgataatc tgaaatctgt 300
 ctagctcaat ggatacaatg gtcctgaaga ggagggcact tcctaccccg 350

<210> 30401
 <211> 485
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30401

ggcatccttt taccttcnga ccgtgaacct agannacnaa cccacgcttg aatgactggg 60
 atgtctgagt atggccagcg actttcattc atctcttgcg gcagcnggaa cagggcgggt 120
 cttagtaata atatgctctc actacagacc aactgtgaa cgggcactnc tattactcat 180
 gatcatgcat ccgagatgag ctcaagggtg tgaatccgca cacagcttgc tcattgtaaa 240
 tgtgtgcaag aatatcttga tccttataaa cgaaacgagc atnattctat aaaagtagag 300
 aatgtatgta gaagtgcctc tgtttataaa tcaccgcatt gaataacaat gaatattcac 360
 ttgttgca taatttaaata gacacacaat atctaaatgt gatgcagtac tcacacgcta 420
 taacataata gntttgtgac cctccccag cgacaatgtt cccgcgacct gacaggatgt 480
 gccct 485

<210> 30402
 <211> 440
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30402

ncccccttga tgcttgatac cacggcgatt tcagctcgca cccgggatcc tcagagtcga 60
cctgcggcat gcaagcnttt tatccaaagc tcattcttggg ggngaagctc cttttttcat 120
ggcttattcc ttatggnatg gcgcctcct ctcaccttta tcctttgtct tccccctgcat 180
ctncatggtg'gaaaatcacc attaaaggac cccatttgaa gctcaaagat ccagcctnca 240
tagaagcccc acaagcaagc ttccatcagt aattntccca gagtgtacag gatagcacct 300
gtccactatc agaaggaaac aacaattaat gtatcaatat atcagcaact aatcatcatc 360
agatacaaac aacaatcaca cccctcaat taattgtaaa gaatacctca aatccttaaa 420
tcaaacaccc tcgatttttg 440

<210> 30403
<211> 432
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30403

tatgtataat ttttgaaggt acagatgtag aacgtttaat tagtttantt aacaaaaagn 60
aaacaacatt tctagcattt gatgagaatg aaatgggata tactacaact tgcacatgtg 120
taaatttatt cataatagta tattggaaaa tcatgagttg cttgtagcat tcaaagaagg 180
agatcaaagc aatattttgc gattatgaaa gccccaaaag tttggggctt agaaatatgt 240
agtgcgatat tgtgattttg tntgagatta gtggaggcta gtgatattga aggttgggca 300
acatatgtgt tacacgagaa attgaaattt aaatataaaa aatagtcgaa aatatganat 360
aaagaacaac tttggaatgt angaaaaggc tgcataaatt gaaagaacaa attatataat 420
ggtattgtag tg 432

<210> 30404
<211> 322
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30404

agctttcntt caacgaagag aagagaaaga cgggagattg cggaaataca agnagaaggg 60
atgtctcctc cacctctagg acctcacaat cactcacaaa ctcactcaa gctctcaaga 120

cggttcctc ttcaagctct ggtctctgct aatcttcaca caacaaaatc tctcaaactc 180
 tttggaactt ggacctttct ctctctataa ctaaagacat gccagagctc ctcaagaaaa 240
 atggccaaac tccatctcta aatctgattt tatgcttaaa taggtggctt tgattgtgct 300
 catgcgctta atgcaactct ga 322

<210> 30405
 <211> 319
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30405

tttgtgctag tgtgcttagc gtgactatgg accgctcagc tcacattagt ggattttggc 60
 ttagtgcgcg ctattctcgc tcaactggatg gactgaagcg gngcgcttag cgggatgacc 120
 cttegtctca tgcaaatgca caactcattc ttgctctaga ttcttcctcg cactcagctg 180
 aggagtgatg cgctcatcgg atggctcgct aagccagaag attggcttat cgagcggatg 240
 aaaatcaaca cttcacgaac ttgcctagat aactttgaaa tgagaggaaa tggttattaa 300
 acacacaaga tgggagttc 319

<210> 30406
 <211> 103
 <212> DNA
 <213> Glycine max

<400> 30406

tcgcgaccaa tttcttgttt gacatcttaa tcttgaattc tggcattcat ccactaatat 60
 cacatatact cgcgaccac catgcgtgag aggtctctatc ccg 103

<210> 30407
 <211> 353
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30407

atggtttctt ctaccctact cctatcgact agtggagcca gtccaagtga tagaggtgtc 60
 atcctctgaa gaggatcctg aggaggacct anaggagtta cctcctgagc ctgctgtgga 120

tgctcttgac cttccagagg atgatgagga cccacttcct gatgtggatt ctccagagga 180
 tatcttgtca gcatttgaga cagactctac agaggagagc ggccctggag ggatagcgaa 240
 cagtgaagac ttttcatcat agcagacgac tccttagact aggcttacat actttttgtg 300
 cgtgggtgta tctaagtcag actgctangg ttactctttt gatttttggg tgg 353

<210> 30408
 <211> 372
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30408

ccacagcaga acaattatga cttttccagc aacagataca accctggatg gaggaatcac 60
 cctaacctca gatggtctag ctctcagcaa caacagcaac ctgctccttc cttccaaaat 120
 gctgctggcc caagcagacc atacattcct ccaccaatcc aacaacagca acaaccccag 180
 aaacagccaa cagttgaggc cctccacaa ccttcctcg aagaacttgt gaggcaaattg 240
 actatgcaga acatgcagtt tcagcaagag accagagcct ncattcagag cttaaccaat 300
 cagatgggac aattggctac ccaattgaat caacaacagt cccagaattc tgacaagctg 360
 cctttctcaag ct 372

<210> 30409
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30409

cgccccgtgg gcttgacnta cagcgaatna ggaacccccg cgggatccaa ggagcgacca 60
 gacggaggca tttttttaag cgcggaaccc ggcgcaacgc ggggtgcatc ccggcatgat 120
 gctcactccc ttgggcgcgc cagtatgaaa tacaagcgac caatgctagg ccggacaccg 180
 ggaatatccg gacataagac atgcaccgtg cctaaggaaa tggcttccca aaagcccaca 240
 agctgagccc aaaaaggacg cccagaaaca agagacccaa cgcaacctgc agcagagaaac 300
 aaaaagaaac gtacatgact ctccaaatgc caatcagaac agaaagacgg ccaacgattg 360
 ccaaaccgtc caaatagcgc gtgctatgaa acaaaaaacg ctggataaaa aaacaccccc 420

agcgaatgg

429

<210> 30410
<211> 511
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30410

nnttatttgg ggaaattccc channngntnn nntntnnnnn ntningcnaa cngnccnnag 60
aggantgaag gacgaggaga ggagacgagc gtttacttag ngttcgacgc cnacnagncn 120
ccaacacagg catggcggct angagcacc naaacacngc aacatccacg ggccacagaa 180
gggacaccgc cgcgagctcc accgccacat tttgacgaca tcgtctttgg agactggaga 240
tacgcaggac aaacacggta tttgaagggc ccatgggtta cagttgccct ctgagaagga 300
gacatgatcc acacgtcagt cttatgggac gacgcccctt attctgacgg tcagacatga 360
acctggatat ctgatcact caactgactg atgcacataa ataccgtaat aaaattcctt 420
caacagcacg tgggacggaa aaagtataga ccgttagcat cgacaatgtc ctcacattgc 480
ggtgatggag ccagaccacc atcgcgtctt g 511

<210> 30411
<211> 308
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30411

agctttttga ccatgcaaac actaaggctt agggttcggt tccccctttc aatcaaccca 60
atgtttccaa aaaatgctcc tttaccaagt catgcataca tccaagtnca tttangcatt 120
tcgggaaaac ctttcattgc gttcaccctt taagcgcaca ttcttttttc ttcaaaaacc 180
tttttgtgtt atgatccggg aattttccaa agaaaactgg cggtcattct ttttaaaaac 240
atgttggcct ttttagtttt ctttccctta gctttttttc ttttcaataa tttctttcaa 300
gcaaaaac 308

<210> 30412
<211> 495

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30412

nnnaaatcga gatcgacngc cttagtacnn nctgnganaa tttagannac tcaccctngt 60
 ccacaaanaa tcaactaaaaa tggnnnattg tctaacttct taaacggnc ctttttgctt 120
 tatgcggtta acatggaccg ttcaaaagca taaaatcaac acatcacttt actacctttc 180
 gcgagaacta cgtangtctg atttcctctt cgatggagga tacataagag caaaaagtcc 240
 ccttttgctg acctgtgag atggttagag gtccaatgcc ttaaattttt tcaccaagta 300
 aaatggatca ttttaaggtc caatgcctta aatgaccacc ttccaagtaa aaagaatcac 360
 ttgattcgcc ccttttgcaa gaactacgta ngctctgatt cctcatcgca attgaggata 420
 cccngagcaa aaaccccgct tttgtcacca cccaagaga actgtatggt ccaaccctta 480
 tcgttctctc ttttn 495

<210> 30413
 <211> 344
 <212> DNA
 <213> Glycine max
 <400> 30413

atatattacc caatttaatc ggacatccga gtaaaaagtt attgtcggtt gaatttccta 60
 cgagcttccg tgttcaattt caacgcctcg atatattaca agactcaacc ggaaattcgc 120
 gtgttaaggt attggcaatt caattttctc agaactttgg atctaaattt tgagcgtctc 180
 gatattattac cggactcaac cagacatctg tgtataaaag tattggcatt tcaatttgct 240
 cacagcttct aatctcacat ttggagcctt ctcatatatt aacccgatcc atcgaccatc 300
 cgagaaaaag aattgtcggt gaaaattcta caaccttccg ttcg 344

<210> 30414
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30414

tctggtggga catcttgact tgctttccaa tcngacattc tctacagatt ctgccttcnn 60

ctatnntcag agggggaatg cctctaacag cacctttgtc aatgattttc ttcacgcctc 120
 ttaagtgcag atgtccaaat atttgatgcc atattttgac ttcaccttct ttggagaata 180
 gacatgtgga ggagtaactg gtttcttgag gtgtccatag gtaacagttg tcctttgatc 240
 tgctgccctt cattaggact tcactcttct catttgtcac caagcattct gactntgtga 300
 agtttacatt gaatccttca tcacacaact gactgatgct gatcaagttc gcagtcagtc 360
 ccttcaccag cagtactttg ttcagactan gaagtccatc atggactagc tttcccattc 420
 cagtgatctt tca 433

<210> 30415
 <211> 151
 <212> DNA
 <213> Glycine max

<400> 30415
 actccatttt tatatattac aattattcat gtctgacatt tgcacgtagg gccctgcaac 60
 tattgttcca ccaatagcta ggaataagct aaccataaca agagccatac caaggaaggt 120
 tgttgattaa gatgatgccg tataaagaaa a 151

<210> 30416
 <211> 425
 <212> DNA
 <213> Glycine max

<400> 30416
 ttgcgactag aatcaccaat agaaccccag cgcaagcgga aagcagaaat tgtacgtgaa 60
 aaaccaccg agcgcgggcg gacgaacacg aacccacccc gagaagaatc ggccaaagga 120
 aaaccggtca tccacagact ggatacctgc gtcacgggaa acagaccttc aacggggaag 180
 aacctcgaac caagcaattc gacacattcc aggcgcccgc agaggccac aaagaggtac 240
 gtgcattgcg agacacagcg ggtagaaccg aaagaccccc cagatgacga gaccaaacca 300
 ggccactaag gcacccttgg caacataaga aaaaaaatt acaccgaacg gtgaacgacc 360
 aacataacac ggaaaaaaa atccgacggc ggcgccgaa acgctcggaa aaaaagacga 420
 aaaac 425

<210> 30417
 <211> 286
 <212> DNA
 <213> Glycine max

<400> 30417

agcttattgt attaagtgaa atagaggatg ataacggtgt gataatccca aggaagtgaa 60
 aagcgaaaca agaagaacca aagaggtaga ggcctgatca accaattatc acatcctctt 120
 ttccaaagaa gagaaagata agacctagat gcaagttctt aaccctaaag gaaccatta 180
 agatcccatc cttcattgtc tcttggtcgt ctgatagtga ggcaactcca actcccaata 240
 cacatccatt cagtccacca ccagtgtaga ccaagaagcc tacata 286

<210> 30418
 <211> 479
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30418

nggcagactt cttcatgcag tatcngacct catnaatact aagcttacct ttttaaggcca 60
 aaccattcac cttgggggtgg caccatgtta tttggttttg caccaanagg tgggaaagga 120
 tgggtccatca tgtgcttgta ggtgtacgat aggtaactca aataacctta ggtaaaaaaa 180
 tgcccttggt tatttggggg tagcaaaaat actttcttgg aaaataatng aatggatgta 240
 tatattgcgt gtagggtagc aaaaatgctc ataatgtat atattgcatg ataggtagcc 300
 aaaaaccttg tggattaatt aactacgtag cacagtaccc tattatttaa gtaattaaat 360
 actttgtggg tttagttaag ataggaacaa aatgcctcta caatgtatta tttttgcan 420
 agaaatgcct cacaaactta tatgtattga tgtaggtagc aaaaccttgc aattaattt 479

<210> 30419
 <211> 344
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30419

ttccaacctt ttttgtgaat tgaggatgga cccctaactt ggggtttgaa taaaaaattt 60
 taaaatttaa ggaaattaaa aatgcctaga attaaatttg tttgatttta atttctttca 120

tttttcaa at gcttttggtg gataaatcaa ttcaaatttc atcaattnta aattcctttgt 180
 ttggataagg caattcaatt ttctccatat gcaaattntc aattttatat tntanataga 240
 tgaaatttta atattaaact ttatagaaaa caaacacatc tattttgaaa tattaattaa 300
 aaatattttc atttttaata tttaaaatac taatattgat attt 344

<210> 30420
 <211> 395
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30420

actaagctgc tgagtgaagc cccagccttg cctggaaccc ttttgctacc tccttccttn 60
 ncccatnccc ctgtagaat tggcattttt aatagtgggt taatgtaagt atgcatgtat 120
 tcattgaact tggagaanaa tagggtaatt aagccgttgc taccatatgg ctttgaaagt 180
 tgaacttaat atgctgtgtc ttggatatat gttgtttggg tgttgcatgt cttcgtaaat 240
 tgaccaa atg ttgtgggtgt gtgctgtctt cttatatatg attccttgagg gatatgagan 300
 aaaagatgga aattttcaatt tcaattacta gtgtcttaga ctctatcatt gagtctatac 360
 tgagagatga cttagtgtat ttatattaat cccat 395

<210> 30421
 <211> 472
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30421

aggggagctg taaccttttg taaaacaang ncgattncga gcncgncccc cgnggannc 60
 ctaaagngga cctgcaggca ggcaaccctt ttttttacac tncaaccang ccnnngcgcg 120
 cgacgagtgc tacagncttc attccttctt tcactttggg tccttccttc ttcttacaca 180
 aattttgttg gtcttcact gatgatgatc atggaaggct aaacactcaa tcaatccaag 240
 gatccactcc aagaaagggt gaatntgagc tctggtttag tatttcaatt acgtgtgaat 300
 gtacatcttt ttcttcaatc atatttttta ttttcatgat tatgaatatg cttaggattg 360
 aaaacaaaat taagctatgg aatcattgtg taatctgaaa tctaatacaca gaatgtttgg 420

acgatattcc aacctaattt ggcacctcaa tgaattaagg attaatcaa cg

472

<210> 30422
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30422

gccgtatann gttttngacn tcnnnnntag nnacnnccaa gnnnnntttt gaagaccccn 60
ttttttgtta aacccaccca ctttccgggg atactcacia atctcccctt gaattgataa 120
agctttctaa agaaattgat actctgtagt cctgaattat cctattcctt ccccttgagg 180
taacaaaagc caaggcgtat agatttgagg atcataataa ctaacgtcat acacattgtg 240
gagaactata accaatcatg aaccggaccg tgagccacat cataatagat atctctatat 300
accataggcg aaacatatta attttgtcca catccatgca atatggaaat taaatgaaaa 360
tccatatatt agccaataca tgctaaatca tgtctag 397

<210> 30423
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30423

aaattatttt gcttgatctt aaagcaattc agcccggacc cgcgagatcc tatagaggac 60
gacctgcgcg cgcgcacacn nttttttcca nagcaccctc tcgagaggag aagccaactt 120
cttatcatag ctccatactc taaatagatg ggtgcctccg tctgatggca aacctactcc 180
catactcggc tattggatat aatgactccc aagtagagat gggacacagc tagaacgaga 240
atgccactaa gggttctcatg agcccttacg agagatttcg ggcccaaggg ctaagtatga 300
gccacttat ctgtggccat acgagatcaa gggttaaata tatctgggcc tcggatttac 360
ggctacatta tgtacgcaag gtacccttg aaagtaagaa acctcaacca cttgatatac 420
ggcccctaga acagaggtcc gatatgggta caacgaacaa tcccg 465

<210> 30424
<211> 484

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30424

ggcgtttggc ttttngttct aanctngtta anaacnangc ggggacaaag gacagggccc 60
cgtcaacaca tttgcttggt gtgcacncnc aannnccggg aggggacgcc gcgnacacac 120
ccncacacac caagnaccan cngcccgnna agggcgncag cccacaaatc cactgaaaca 180
aagactcgag cggccatctc anggatacca tgtgatatga gaaatgcctt tttcgtgggt 240
agtgttccta cgtaacagtt gaacctagtt ctgcagccct ccatatttaa cacacttatt 300
gagaagatct accgtctgct catcgttggg aaaggatgat aggagaagtc atcaattcac 360
atcctgtaat ctgatttggt aataaaaacta ataaattact atattctaac taatacaaca 420
atattattcg agattatcat gcttctctct tgcggttcaa tggcataaaa tccgcttgct 480
cccg 484

<210> 30425
<211> 324
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30425

agctttttat ctttcattgg tgtantttta tccgcttttg gtgctctaaa ttgtgggaat 60
gtgctcanat atgtggtgca attttggttt gttttcttgc ttgattgggt tgaattgngg 120
gtttgtatga gatggccta tgcctataat gcattttgaa gcaatgggac atgccacatt 180
gtccccgttc tcttgcattt aatgcctaaa cgcgcgccca ccaagtgttc ggtgaaatgc 240
ctcaatggca ttaacgcgtg attttcctan ggaaacaacc catggggcat tttggtttcc 300
acatattttc tatttttttg gaca 324

<210> 30426
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30426

accccctgtg ccctccgttt gatttaagcc aagcccctac ttttgagggg caactcctac 60
 cttatgaaga ctatcccggt caagacgatg gngaaggaga tacccatctt ggccccctgc 120
 tccacctcan agatccatcc ccgcatgaac taccacagct gaacatagtc cgccatatcc 180
 cggcctcatc cgcacccgta aaagaatcta ttccctttgc ggaagataag ggaaagattg 240
 aagcgctcga agagagggtta agagcagtcg agggcctcgg taattaccca ttctcagatt 300
 tggcagatnt atgtcttatg cccaacatcg tcaccccttc caaattcaaa gtactagact 360
 nntgatagt 369

<210> 30427
 <211> 297
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30427

agctttttaga ggatgcttta atgaaggana agaaagagag atagtgggag cacgaaatgg 60
 aaggaatata agagggagag aagtgggaact ttgaagtgtc tcataagact ttcattcatc 120
 aaagttacaa caagtgttac acatgcttct atttatagac tangtagctt ccttgagaag 180
 ctntcttgag aaaacttcct ttataagcta aagcttagct acacacacnc cctcttaaag 240
 ctaagctcac cttcttgaga agcttncttg agaaactaga gcttagctac acacacc 297

<210> 30428
 <211> 510
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30428

nnntttactt ttngcttgat antgcggcca ctctatagan taccctcgct nggtgaaatg 60
 ccacattcat ggtcattgtc ggagnaaatg catganttat aaatcttggc catatgcnca 120
 gacgcgtgtc tcgtagaagc atatccattg ggatataatg atgaaactat gtgcattntt 180
 caggtagaga aagacggcta gagttttgaa ttgccacaaa gtagcaggtt ccggctaagc 240
 gcatatacat cactatgcgc gagatcagtg cgctaagcgc aggatgtgcc ttcagccaat 300
 gtaagctcga gactggcgct aagcccaatt tcacttactt gcgctgagcg ngagggtggc 360

gcttagcgca gcgtcacgag ttcagagcct atntaaagcc tgtcttgtgc agaatagggt 420
 acacaccttt tatgtcatct tctacacact tgtcacgacg accagggcac agaattcata 480
 gcccgcatatc ggctatttgg agaaaaagcg 510

<210> 30429
 <211> 173
 <212> DNA
 <213> Glycine max

<400> 30429

agctttttgtt agatgccccca gctaccggaa aaagcggcct tgacgactcc aagagaggct 60
 atagagacat tgaaaatccg acctctaata ctgatgaatt tctaaatgat agaattcatgg 120
 ctccgtcttg aagactattg acagtccata ctcaagatgg tctcttagcc caa 173

<210> 30430
 <211> 459
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30430

nccttaccag ggttagnagc tgtgctgcgc ctctatagan taccctgcct gagactgggg 60
 cgaaggccca ttgccgttac cncngcttat tcaacttttcg gagaccaaaa ccctattgag 120
 agtctatctt gtgcagaatt aggggaaccac ctttaccact tttatgacaa cttctacaga 180
 caaccagggc ccagaagttt gaaagcagcc accggcctat tgggggaaaa gagccctaga 240
 agcagatata tgagcagcct gtgcattgaa gcctacgttt tgcattctga aaaaatattg 300
 gtagagagga ctgtatatgc tgataaagga ggggaatccc cttcttggaaggactatca 360
 tctttgtttt atctattatt gtaagggttt tgtatggtcg ctaacaccct atgacgattg 420
 ctatgacact aatgaaacct atatctatga tgggtcatg 459

<210> 30431
 <211> 361
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30431

agctgtgtta ctcccagaag actgnaagng tgctactggt gcaatttggt ttaactgtta 60
atcaatttgt gttttatgcg ttgataaata aatataaata atggataaca tttctcaatg 120
atctttagt gcccttggtt tgctgtgtag ctttattatc cactggctag tacaaaaaca 180
ccatcttgca tgcattgctt gatgggcgt ggcaacatta attttataaa cctatttcta 240
gttaaaatta attctaaagt gatatgatgt atatttaa atttttatta taaaactaag 300
aagctaagt tataaaactc aattaattct ggacgtacac aatcaatttt aaactctttt 360
a 361

<210> 30432
<211> 228
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30432

tttttataaa aaaaaaacgc gaaaaggaat tataaaaggg aaatgaaaga tataaattaa 60
tagagaaagt tgggtggttg ccattagaat taaaaggga ttttgagat tntattttat 120
tttttggaat ttgaaaaaat taaaaaacat tgttaaaaaa aaaagttaac gaaatcaaat 180
gataatattt acggagaggg aatgcgattt tttgatatag ccttatct 228

<210> 30433
<211> 403
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30433

agggagtacg atgtatctcc gnatttataaa gncccgggac ttaagggacc tctggaggcc 60
cgtttgatcc anccgaaaaa acgggcgtgt atgagaaact cgacataccc acacggcaag 120
caaccatgaa tggcaacatg ggtcccagaa tatacttgaa agccggtgga tagagtccac 180
cgcacataca gccagggaa gcttactaac aagccacgct atgacacgaa gcaatggtat 240
ttatgacaaa acaccctgcc tctactatgt tagcgaaatt cctgatgtcc ttatcgcaag 300
atgatccgta aggtacagcg caagagacat taggtttcct aatacacaat aagtgggagc 360
gccctcaatt cgtaaggggag acagttggtg gcacattatt tcc 403

<210> 30434
 <211> 490
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30434

 nttcttttnn ggatccccnn gntgcntcgn gtnanttana cnaccncact cttgcgagcg 60
 agctgttgag acacacacaa ccggttagtc agtttttgac accccaggcg ccaggaggag 120
 agcgcaacac caaaggagac aagacacacn caggcccgac gaaggggaag gagatagcgc 180
 tcgcggcacc cgggtccacg agcacaaaga gcggccgggc gaggagcaca gaagaacaga 240
 gccgcgcaa aactcgacgc tccgcccccg aagaaaatc catcgcgga gacacaccac 300
 agggaggggac tcacgcggat cgccagacga tagtaacaac tgtgacgac gccctaacca 360
 tgatgacaaa cgtecccccg agatcaagtc ccatcaccat cggaacgaca aggaatcagc 420
 tgtgaacaac tggacctaca aatggccgac atgaaacata cagctctata cgatgagaga 480
 cacatgcggg 490

<210> 30435
 <211> 238
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30435

 agcttttagc caaaatcctg actcaccata aaccttgacc cagggtgaga atgtcaatcc 60
 ttaccctcgg aagcgaanag aatagaaggg aaatttccaa tcaaagaaaa ggaaagaagg 120
 aagatttcca atcaaagaga aagccaaaaa agaaaagatt gaaaattccc aatcaaagag 180
 tgggagaaag caaaaagata agaaagaaaa ttcccaatca aagaatggga gaaagtaa 238

<210> 30436
 <211> 220
 <212> DNA
 <213> Glycine max

 <400> 30436

 tgccaactca tatgagggat aaacacataa gatcagtcct gagaaaaaat gtatcattac 60

tgatacacac caatccatgt catgaaaaga aatgtatagc gatatactat caaatatgat 120
 tgcaccccaa tttatacaaa tgggtgtgtt tcttttgata tatgaaagaa actttgatca 180
 cgctttctgg atctacaatc agatggacgg tataaaactta 220

<210> 30437
 <211> 316
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30437

agcttggttat tatccaaatt ctgagagata ttactcacca tgagaatgag atcactntta 60
 ataacttcat cttgggttaat ttcacatca atttgtaata ttgatcccat aatatactta 120
 aaaactgggtg gtatgatatc cacccaacag tcacacaatg tttttttact agtaattctg 180
 tcaatgatat gagcaattnt tcttattaga atagaccttg actctactat tatagcgaaa 240
 tttctgtaat ttcttttcaa aatatgaacc taaattaaaa gagaaagaaa aaataaatat 300
 atttattata caaaaa 316

<210> 30438
 <211> 424
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30438

tgttcgcaca tcgttcgcgt gtatgatatt cactccacaa ttttttaata taagaaaacc 60
 ttcaatccta taacgcacgt ggcgacaaaa tgggcataac tgaatggcat tattgcaatg 120
 cggaagggtat tctgcgcttc actatccatg ttcacacatt atngcagctt gtggttacgt 180
 gagcatgaac tactaccaat atatagatgt tgtttacacg aatgagcaca tcttaaaaagc 240
 atactccgca cagtgggtggc ctcttgggaa tgaagcggca attcctcctt ctgatgaggc 300
 atggacatta atccctgacc caactacaat tcgtgcaaaa ggtcggccaa aatcaacaag 360
 gataaggaat gagatggatt gngtcgaacc atctgaccac cgacanaaat gtagtagatg 420
 tgga 424

<210> 30439

<211> 350
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30439

accttatcta ggttggacct cggctggagt tgaatacgta aggctggagt ttggctcatt 60
 gcctgtcata ggtttttntt aaagctcggc tcggtttaca taaaagtctg gctttgcccc 120
 cgagcctatt taaaaacttg cttaaagacg tctttgatta attaattatt ttaaaatcta 180
 gtgaaatact aacttaaaaa agaaacttat aaaatttcgt ataagcaatg taaaaattca 240
 aaaataattg gataacaaaa tcatattgaa ttcaagtcgt taaagtacaa agtatatcaa 300
 aagaaaataa aaagagcata atattaataa atgtatggat tagagatgat 350

<210> 30440
 <211> 233
 <212> DNA
 <213> Glycine max

<400> 30440

gactgatcgt tgccctttct ctgcgctaaa caaacagaga acgtcgctgc aagacagccc 60
 cgtatccttt gtattcgcag gtttctttta ctaatttggt ggcttaaaaa gaaaattata 120
 ataaataata agtcgacgcc taaattctaa cttaagtaag ttcaagttag gcaagacgct 180
 aacccatgag aaaggagggg acatgggttaa tgttcccctt cagaaaaaaaa aat 233

<210> 30441
 <211> 286
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30441

aataactatt ttaaaatgat aatttagaca aaaaattaac aaaatcattt ggattatctc 60
 ccaatctcac ctaatacata cttcaggcgg tagctgcgtt gggatggact gtgaatatat 120
 agtccgtgcc tgcgtaaata agattgcccg tggccttcct ttagctacgg gcganacagt 180
 tttatggtgt taacctttct attatcccat cccaaatgct tagacattta agacaagccg 240
 atctacatat taggaaaata acaaagtca tcatcataaa aaaaaa 286

<210> 30442
 <211> 439
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30442

 nggagtgtgt tcatgagaac gcaccaana aactcagcct ngctgtgagt cccagaaga 60
 ttttaattggc gaccatttat tattttttaa ggaccataaa aaatgtagga gtctatcttt 120
 caatcttctc tcaacatcat tcaatatctt tcaactgttt ctacaaaaat atcttgaatc 180
 attctctctc atcttcctc aagtcttggt tcaacacttt cttttccaaa acaagtcttt 240
 gtcaaaaact cgtgctatca tatttttcatt ctogtcttct ttcccaaaga caaagactaa 300
 ccgctgattc tttgtgctct ctcttcttac aaagatcaag gacaaccgct gaaatctttg 360
 ttcttctctc cctagcaaag attcaaagct aaccgctgaa tactttgttc cttacaaga 420
 ttcaaggata ccgctgaaa 439

<210> 30443
 <211> 365
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30443

 agctttgttc ttgacanaaa ataacatttt gaatggtgtt aatatatata ccagtttact 60
 aatgtatata tacttgtttt ttttaatatg agtacgttaa caaattatac ctagatatta 120
 tcttacaata aaccaataat ataacttatt aacacactta aaatacacca ataataacc 180
 tagatatttt aattaatata taaatagagt tatattatta ttattaataa atactccaat 240
 atttctatga taaaagcaca tgcactttga taatgaaaaa ttacctttct tataaaatat 300
 gatggttcta tgaatctagc atactatgaa taatatataa agttttttta ttcataaata 360
 tcaaa 365

<210> 30444
 <211> 442
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30444

aacatctgga ccttaggtgc cattttttaga gtaaaatttt aagttaattt tgggggatna 60
aaaccattct tagagtccaa attaaaaatt aaaaattcaa atcacaattt gtggttgtag 120
acaaccacca cacacgttcc agggcaaaaa attcaaattg agggcccat tgtggcttaa 180
acaacgggtg tgcgtggcaa attcaatggt atcactgaca acccatcata gtccccacg 240
catcttcaag agctttcaat anggacanat ttangtcca aattgcaagt ccacccccgc 300
aaagccctca ccctacaccc tcacagatct ggttgatgat gtagaggcat ccacggagg 360
aaaggcagtg gtgcacaatg tcanggttga gccataaag gatgtggaag gccgacttga 420
tgaagaggaa gtcacgtca at 442

<210> 30445
<211> 222
<212> DNA
<213> Glycine max

<400> 30445

agcttttaat ttaatacgg acatcaaac agagcagtgc attaccaatt taagtactac 60
ccaccaccaa gcaaagctat gttgaagata tacttttgta actacataat attttatttt 120
ttcagttttt acaataggat ttagtaaata agttggtgtc ctatatTTTA agagtaagtc 180
agttctaatt gattagctta gtcaaaaagt ggttcctatc tt 222

<210> 30446
<211> 209
<212> DNA
<213> Glycine max

<400> 30446

ggcccgtat gtaactaggt tctatctact gcaactgcta ttatcccaa tctttattgg 60
attttatata agcaaatgaa gtgtgaggaa aagtaaaaat tggatcataa agaagaaaaa 120
attgtgaatt agttgtacat acttttgaat tttgcactat ttacgagtac ttaaagacaa 180
tattacttat ataatgatta tctaaacag 209

<210> 30447
<211> 179

<212> DNA
 <213> Glycine max
 <400> 30447

ctgagtaggc tgctgtatga gatctctcag aaggactgaa tgcttgggca cagaatctac 60
 taccatagtg cttgggcaaa gtggaaaccc ccaatccacg gatttggata tgcagcttag 120
 tggacaagga tagacaatat agctactttc ttttttaata tacctatgtc attattgct 179

<210> 30448
 <211> 493
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30448

naacaggaat tttagtnant gcgaactata gaaacacaac ccgggggggat atcactcgcg 60
 ggtacgaacg aaccgcagcg atttgcattt cattcgccac ccactgacga gagcggggtg 120
 atgaggccaa aagcctgaga gctacggagc ggcccctgcc gtagacacag aaagcaaccc 180
 ttggagttgc tgatgctgag acaagagcag caactcccac gtcactggaa gcagcactcg 240
 agcctctaaa ctcagcatga ccagacaaaa cgacgcgcgt caagaccagg agaagagaac 300
 ctggagcagt gtcatectca gtgagacaag acgaagggga ggtgctgccc gttataagag 360
 cagagatgc ctaacgaaac gagacattag aagccacaat gccgacagcg ggggaatgat 420
 attcatgtgc caagaccaag ttcggagtca acatgtctgg atggaccagc tacgagataa 480
 cttccctccg cgn 493

<210> 30449
 <211> 303
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30449

agcttatatg aacaaaattg ccttaatcat tccaaatatg catgtgaatt angacgcac 60
 aacaagaatc aagccaaggc tattgtgcaa gcaatcaatg gggcaaaaca cacccaatga 120
 ttataatgat ggatggctca nattctcaca aaggtaaaat catcactttc aaattgagct 180
 ttcaaaacta tcatgacatg tagagaagaa tcaaggattt caagtcacaa aatgtcaaga 240

actttttat tcaaaacaat tacccatttc ttgaacatat cctataattc aaagaaaaac 300
atg 303

<210> 30450
<211> 134
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30450

ntgtcctcag atccattgg tgggactagg ctcaatttag tcggttctcc tatgnntaga 60
ctaacttana ctaagcttca tctcagatc ccatttggtg gactagactt agcttaaata 120
gcttatgaaa tttt 134

<210> 30451
<211> 244
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30451

agctngtgaa ccgatataatc gataatattt tagatcgaaa gatccttccg gaaaggaatg 60
taaaacttta tcattctgaa tttgacgagt ttaaaataga attagagagg cgaaacctgc 120
acaaacgtct cgccaacctt caggaaggaa gcatagatgt ggcagtgggt aaggaatttt 180
atgccaatat ctatagtcca gcaaataag ctcttanata tgctaaaaca agaagccatt 240
taat 244

<210> 30452
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30452

ggctcaacta caaccttattc gagggcctca aacactattc gtgntccca ctccagctn 60
tgtttactat tccatcgacc gacagaacca aggggatgaa ctccaatctt cggtcgacgg 120
aggagtgagc tctacaacan agtcaatgag cacctagtcc ttgatggcac ctcttttttc 180

aaagatgatg ccatacttgg acaattcgat caaccacttc accatccttc ctcccaaata 240
 gggtttttac agaatttttc agattggctg atccatttgg atgactactg ggaaactcta 300
 aaagtaatgt tgcaatctct ggacgtgatt agcattgcc aagaccattnt gtctaactct 360
 tgataatgcg actcaacacc ccacaaaaca tggctgatga agatgttgaa gaaatagagc 420
 agcacactca gt 432

<210> 30453
 <211> 240
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30453

agcttttttt accatgagat tgtttgaggc cttatgtttt tcttgatctt gtntacttga 60
 ccttaaatac atgttgaagc aatgcttaac ctttgaatgt atggtgaact aaccttgtat 120
 taatcttaaa gcaatgctta acctttgaat gcttggtgaa ccaaccttgt atgaacctac 180
 attggcatca tcagaaccct gtatacatat attcacaata ggtacccgac tacgtgtatt 240

<210> 30454
 <211> 426
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30454

gggccgctgc cattacgacn acnnncctta tnganacnca agccancngc acaaagggca 60
 cggatggcac ccaaggggaat gattaattca gccccaaac caaacngagg ggacaacaca 120
 annacaaaac aaggaccaca tgccccgcac gcaaacaag acagcgcta cgcaaagaaa 180
 atagcccgcc acggaaagag agacacacaa tgcgctggca gagccaggag gcaaggacca 240
 accaccgacc ccccgccact agaccaaaaa tatcaagggc acaagcaagg ccggccacga 300
 agaacgcccc ggataacgcc ccgctcgagg cacaggagcc caagaaaaca gctacaacca 360
 aacgcgggct ccttggccta agaggccaaa gaagaagccc cgccaaaaac caggacacgc 420
 tcgacn 426

<210> 30455

<211> 376
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30455

```
agctttttatt gttcaatttc gagtgtctcg atagaggatg cccctgaatc ggacctccga 60
atgaaaagtt atgaccattt gaatttctcg agagctacct ttgttcaatn tcgtgcgtct 120
cgatatatta tgcgcctgaa tcggacctcc gagtgaaaag ttatgaccat ttgaatttct 180
cgagagcttc cgatgttcaa tttcgagcgt cttgatatac tatgcgactg aatctaacct 240
ccgtgtgaaa agttatgacc atttgaattt ctcgagagcc tccggtgttc aattttgagc 300
ggctctaact gtgatgcgcc tgaatcagac atccgagtga aaagtatgga ccattgattt 360
ctcgagagct cccgtg 376
```

<210> 30456
 <211> 349
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30456

```
tccttgtggt cggactctca gccacttatg atagccgtcg atgatcccat tactgcttcc 60
cctaagetct ctgtcctttc ttcacgccgc atcccatgcc ttgcgaactc cttggagtac 120
cctcgcgttg tggctactga naccctgtgc gatgaaaggc gtgatgcttt cgtctaattg 180
cgctcctctc atggggtagc caagctgtct tatggcgaga acgggattat aattaatata 240
accccttggt cccatcaagg gaacatttgg acatccttcg catgaagata gaatcttgat 300
tcttccttcc ttctagcgag ggaaccaatt aacagaacgc ccccatgc 349
```

<210> 30457
 <211> 206
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30457

```
agttttatatt tctgattgng gacctgtggt ttgtgcaagc gcgtcaaaag tctacgcacc 60
ttgaaatggt cttgatggat gcaaaggat gttgcgattt agcttttgct ttgtttaata 120
```

atgagatacg gattatgctc tgctttgctg attgggtggt ttgatcccc atattgagtt 180
gtaatttatg ggatttggtt aatcat 206

<210> 30458
<211> 510
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30458

nnggggcgga gggaganann nnnnnnggat ccccnngna tnanactnng nanaananc 60
nnacnnancc nntnanattg cnacatgnng aggttaggac acagtttgca tancettaata 120
tcgcgnangc aaactgggag tctggtgcat actgaaagcc catgaggccc actaaacaaa 180
tctaagatag ctgatgaatg tgtatactaa tgaatccaac gctgggacgt cagatgacaa 240
tggatacacg atggtcagaa agacaaaatt tggccacata tatgttaaga ccatgtcctg 300
cgcacaccct attgaagagg aggttacaag actgcgatct tcaggttcta aggcaagaac 360
gcatttccca gagccataca tacctaaaca ggctctccta tggatgatcc taccatgagc 420
ctgcggaatt tgagcacaga nagacacatt catcacaata cacgcactct gttagatttt 480
aaagtgggtgc gaccttcggt cattgaaatg 510

<210> 30459
<211> 493
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30459

gaagttcctt gattcccaat tgtanctccg ggntaaagag atttgagaag gtttgtgatt 60
tgaacacaan gccacggggg gttttatata attatcacct cactttaatg ctgctatgct 120
cataagtaat tgatatgttc aacggtataa ttgtatacgt tgttcatacg ttacatctct 180
agtgtatcaa ccgcctcggt gataatatta gagcatggat gattgggtata ttgctgttat 240
atatattgct ttaccgcttt tcggtgtcat ggctaaatta ttaccttcg ttgctaaaatg 300
cgtaatccgt acagacgatg atctacctct aatgctatta tcatgactcg taactaatat 360
tatgaatact atatctcttc gtgtttttca tagatgttca ttcattactt acagtacttc 420

tcgtcttaca atacgaattt atagccgtac tcgtgtttat agtatagttt atacacttct 480
actcatctac tcg 493

<210> 30460
<211> 389
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30460

ggggcgtgaa tcatgacatg gacnccnenn nnntaganaa anccccggag nangaacgaa 60
agagggggga cgaaggatat ttcccccca cgccaaacgg ggggggaggg aaaccaaacc 120
acccacgac aaacacgcg agacaagacg gaaccagaaa aaaaagagaa cggggcgaaac 180
cagccgacgc acccagcgaa gcgcaaaaag acaaacgcga ggcccgacac aagagacgca 240
gcagacccag ggggcgacac ggaagagggg cacaacaaga gcaaaaccca gcacaccgga 300
gaccggagcc cagcaaaagg ccgggacaac gcagcgaacc acgcccacga agacagcaca 360
agacccgca caggaagcgc aagcaggcg 389

<210> 30461
<211> 297
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30461

ctcatcttgg gggagaagct gcttctttca tggcttattc cttaatggat ggcgcctnct 60
ctcacctnct ttctttgtc ttccgctgca tcttcatggt ggaaaatcac cattaaagga 120
ccccattgaa gctcaaagat ccagcctcca tataagctcc acaagcaagc ttccatcaag 180
tggtaatcag agcacaagag cttcaagtag gtgctcctta aacctccatt aattntttt 240
ctttaccttc tcttcattg ttggttcttc atttttatcc atgtatctcc tcacatg 297

<210> 30462
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30462

cggcgcaagc gctgagacct ctgactagng gcngatagtt tattttatcg cacatcacta 60
cgctggggcg atcatcttta ctggaagtaa acttcaagca gtgggcttag tggagatgat 120
tgttagtcaa tgaatacgac taacttttgt gtaagatata tgtgaaaatt gtatctaact 180
cctcccattt atgggttattt ggtagtgttg taattacctt ttgttaaata taggtcataa 240
gtacttagta ctcccatttt gtgtatttaa taatcatttc ctttcaattt cagggttaatt 300
aggcaagttt gtgaagtgtc gaatttgata tgctcgctaa gccaatctgt cggcttagcg 360
agccatcccc tgagcgcacc acatttgtgg attatcgcta gacagaatct tgaagaagga 420
tgagcttgac cactcgct 438

<210> 30463

<211> 216

<212> DNA

<213> Glycine max

<400> 30463

atacacttcc ttcaaagtga agtgtgtagc ctttctccat catttggcca atgcttagaa 60
gattttcttt taggttgga actagtaaga caacatggat gaatcgctta cttttatctg 120
tctccaccgt tacagtgcct atgcctattg attctaccac actttcattt tccagtcgaa 180
cttttacttt tgacaacttg gcaatgcctt tgaaaa 216

<210> 30464

<211> 477

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30464

ggcagaatga tccgtcagat cnancnannn nanaannnac ctncgtagt ctttcatctt 60
gtttttaaac acgaaaggag ggtantttta ttttgatnct anggnccgaa gtgagggaga 120
gaagcttaat aaagttactt gacaagagag gcttattaaa gtggaatttc aaaattgatt 180
cgaaaaacca cacctggcct tcaccaacct taagttattc cttgacaata gatgctgtga 240
aatatcatat tgttcgcgaa tttcaggaaa ccctaattgt ttcaaaaagg cgaaaactgc 300
cattaaacta ctaaagaaca tgaaaggcct tgaggaaatg ttcacatttc aaaaagcgac 360

ccttatggag ggtttcaaca tatattgaaa ttgatactg aggacaatgt ctcaggaggt 420
acgaatccgc ttagacatgc gaatgttatg aagttgagcg ataaacagta atacacc 477

<210> 30465
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30465

aggacgtcga tcacttgtat ncccccaat nttaggataa tcnccgggat cctatatagt 60
cgaccatgaa ggaggcaagc gttttgttct ctttatctaa cactaatcct aacgggggct 120
taaatccctt aaattcctaa gctacagcta agtcttctac ccaaaagtta agatagaaaa 180
agagaaaaag gatcaaggaa cttacttggga ccgtgtatga acgatgcttc aaagtccaaa 240
aaggcccaaa gagagttcaa atgcatgatg tgcaaatttc tttggagaga aagaatgcac 300
atgcgaagtt tctgtactat aacaaatttg agaggaactg gtgggttact cactttaaca 360
cgttggaact ttccgttaac gggacatttc gctaatagagc aaaaaatact attggttcta 420
aaccaacttg cttacgaaca gggctn 446

<210> 30466
<211> 254
<212> DNA
<213> Glycine max

<400> 30466

catatgctga caatagccga gaaacccatg aatctcttct ggggcggaga aggtgtctgc 60
catcgcttg gccttgcta acaatcgggg aagttcttga ctcccggtca atgaaaagca 120
caccgatcca tccacatggt tgctctttg tgtaaagagt cgatccccct cctctaccct 180
cttttccgct atacttgtgc atattcgtcc gcatactatg ctctggccc gcggtagacc 240
ctactctctg gtac 254

<210> 30467
<211> 286
<212> DNA
<213> Glycine max

<400> 30467

aaatcctgac tcaccataaa ccttgaccca ggggtgagaat gtcaatcctt accctcggaa 60
gcaaaaaaaaa gaatagaagg gaaatttcca atcaaagaaa aagagaagga aaattttccaa 120
tgaaagcaaa ctaaagaata gaaggaaaat tccccaatca aagagtggga gaaagcacia 180
agaaaagaaa ggaaattccc aatcaaagaa tgggaaaaag tttaaaagga agaagaataa 240
ggaaagaaaag ctctgatca tggatcgaag gaaaaacaga aaaaat 286

<210> 30468

<211> 487

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30468

agaagtttag ttcatgnaga cgnacacta tananactaa gcttctacct cgaaggncca 60
atccagccgc atataatata gagaaccttc taaattaacc aacggaagct ctcgagagat 120
cagatggcat anactttacc tcggaggctg attcaggcgc ataataattc agaccctaaa 180
ttgaacaagg aaagctctca aaaattccaa atgggtcataa cttttcacac ggatgtctga 240
ttcaagcgca taatatatcg agacgctcca aatttaacaa cgcaagctc ctcgagaaat 300
accaatggtc ataacttttc actgggatgt cggattcacg cgcataatac attgagacgc 360
ctcaaattga acaacggaag ctctccaaaa attcaaaggt cataactttt cactcggatg 420
tccgattcac gcgcatcata tatcgagacg ctccgaaatg accacggagc ctctcgagat 480
attcaat 487

<210> 30469

<211> 386

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30469

tantgcacia ccncaagcgn cgcccaaggg cggggctttc agaancagga cctcccttcc 60
aatatgacca ggaccagagc cattaccttc gagatgacaa ttggacttgc tcatttcctt 120
tcagagaaat tgaaccact tataattgac cacagatgat acattgagaa gtcattagaa 180

tgggaataag cactgcataa taaaacttca cactagtatt ttgggacata aagcacaggc 240
 atacatatga ttaattcaga taacatccaa tgtttattga tgtcctcctt tgggtgatca 300
 cccacacaca acatatgaac atgatgatgc taataaaaat cttgacatta tgtgggcaat 360
 taatatgccc taactgtagg tgctan 386

<210> 30470
 <211> 490
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30470

gcgatgctat ttctttatnn ctnncannna naganacnch gcgcagacca ggcaggnggg 60
 aataacgagc ccccccatg atttagttcc acgccccan naaaaaaagg cgcgggagac 120
 cacggggaca cccacacaaa gacccccaan gggacacact accnggaaag acgccncgga 180
 cccgccaaag aaccaacan gggaggacac ncacgcaagc gggagccaag aaacaagcgc 240
 gggaaaaagc gcgcgaccaa caaagcggca agaacnggcg cgccaacgca caggaaggcc 300
 accgaaacag anacgggatc aaagggaaac ngacagcccc aaggaccaac cacacagaag 360
 acggcgangn caacnganc cacgagagg aggccactcc agncgccccg cacacagaga 420
 aaagacaacg ccgacaacga cggcgaggan gaagcccatt cccccgcaca gcaagccggg 480
 aagacgagcg 490

<210> 30471
 <211> 347
 <212> DNA
 <213> Glycine max

<400> 30471

agctttatgc cctcaagcag cgcttttcac atgctgcacc attgttccgt gatagcatgc 60
 acactttccg caccaacttt ttcaatgtta ttgagtataa gtagcatccc atctgttttt 120
 ttttcatgat gtcaatgaaa tgaatatgca tggcatattg acatcagcta atttataggc 180
 tcaaagaaag taggaggagg aaaaccaatc aataaatcat ttttggagg ctgaatttca 240
 cccaaagaag gacctattgt tattattaaa aacagaccag accttactca cttgccaaag 300
 ctagctcaat tgacatctta atacaccccc taacccaaaa tggacat 347

<210> 30472
 <211> 483
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30472

 nggacttagg ggcttgatnt ctannnatng anannnacna ccgtngaata tntggataat 60
 tctgacagga cagatttat tatattgca ctaatgatna naggagagcg acccaatgag 120
 aagccgacac tgacggaaaa tcagaggaag tccctaatta aacctaaaac aaggaaacaa 180
 gtgagcaagt ctttttttct tagtgtgagg gatcaacacg caacnttttc tcttatatat 240
 gtctttctta acccctcaac aaattgtata tcttttaggt tattgaaaat tgtaatagaa 300
 cattaagagt atattgtttt tacaaaatag aaaaatatat tttagcttgc ataaataatg 360
 ggaaacttta tagtaaaaat ataatacttt gaggatataa attctggtgg aactctatat 420
 atatatatat atatctatat gtatatatag atatatgtat atatatatat atgtatctat 480
 att 483

<210> 30473
 <211> 367
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30473

 agcttttgtt aatcgattaa acatatttgg taatcgatta ccagtgtttg tttctgaaaa 60
 atctaaagat gtaactcttc anaaaggttt tgactttntc aaatgggttt taagtttttc 120
 taaaaagtta taactcttct gaatggtctt cttgatcaga catgaagagt ctataaaagc 180
 aaggctntgt tttgcatttt gaatcaatca tttttccaat ctttctaaca aactcataca 240
 atcttttaca agccttgaat ctcttgaaat ttctttgaac ttctttcttct tctttgtacc 300
 aaaaactttc tgaagttttc tggttttcca aaccttgaaa acttgtgcta ttcattcttt 360
 tcattct 367

<210> 30474
 <211> 346

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30474

 ggggtannaac gaaatgttgt caaatacaat aaaagtataa tttgtttata ttntnnctta 60
 nacaatggag gatgtcttca agcgtgatcc gtgtgccgcc ttcacgtga cctgacataa 120
 acattgcact ctacattgac acccctagtg atccttaacc agtcttgcac gacacgtgt 180
 gctttcgtgc cttcagtcac tatcctgagg ttgagcaacc actccaacct ttctgtaatt 240
 gcttggcaag cctcctatga cattgacaac aacagataag gtattaccat attgcataat 300
 taaattaaat gaaattaaag agtacatgaa tttatatgtt accact 346

<210> 30475
 <211> 318
 <212> DNA
 <213> Glycine max

 <400> 30475

 aggetgagac tttggtgacc caaacacgct atggggggga gctctcctct aacttgccgc 60
 acttgtttca tcgctaataa tcaataagaa tctcatcact taattattta acgtccctga 120
 gcattaaaaa tattccgaaa cattgaatta cgctcttttt ataatcatct ctttaaaaac 180
 tttggaactt agagacagat ttaaaataaa attggaaacc tgaaaatatt tttattactg 240
 aattttacta cttaaattta aagggttttt taaaaatcaa ttccatttct aaacatatgt 300
 tgaaacgatc aattaaat 318

<210> 30476
 <211> 374
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30476

 gggtcaccca cgacccccc nataaaccac cgaccgaaac gagcgagaga aggaagcata 60
 tttatcgagc cagcccacca gcgagggggg gccaggaaac aggaccaca ccccgangcg 120
 gaaagaacca ccgcagaacg caacagaaag aacgcgcggc accaagaaca cgcagaaaca 180
 cccgggaaag accaccgcac ncgacggaag ccaaagacac caaccccccc aaagaaaagc 240

aacccaancc cgaacggcca cgcagcaaaa caagagacca cccacacaa aggccacgga 300
gaaggcgggg aaaacaaacc aagagacgag cgcccacgga cggaaaaaga ccgccacgga 360
aacagcaggg gacg 374

<210> 30477
<211> 250
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30477

agctngtgng ggtcgtgggc agactcaaaa ataatggaat gtgtagtggc tttgtcccat 60
acctccattg cagatcagtt gtgcctgagc ttgcttcctt atgtttatta attttgttca 120
tgtttatgaa ggaatcgaaa ctttctgccc aggaccataa cgttaaaaaa ttatgcatat 180
atgaacaaa ataatgtttt aaaactatag ggactataaa gaaaaattat cacaactatc 240
aggacttaaa 250

<210> 30478
<211> 441
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30478

aggagacca tttcatgcga tcgtacnnac nnnttagaat agccttccta ctaccagcta 60
ataattaacg atgcactaag accattcttt acttcttaan gcgngnanga atcgaggatc 120
agtatactaa gccagactac aatgtaacct ggctattatt catactgtgg ttctaaaaaa 180
agaaagaaga gtagatcgcc ttgcttcata taaaagaaag taaaagaac actgtcctct 240
gtatttgtgt ttttcaatac aaaagaagaa gataccctga gaaaactgat cctcctcagt 300
cacctttttt ctaccacatt aattaattgg agcaacaagc tgatttcttc tccacacaaa 360
cagaccactc cctcagggga ttatgtttac cccacaaca taatgcactg cagttaatag 420
gggacataat aatgtttttc t 441

<210> 30479
<211> 289

<212> DNA
<213> Glycine max

<400> 30479

atggcatgat gcagatatca ccacgtactc aactctgata aggacactta attgtgcctc 60
ttcatgcctt agtttgatgc acttggcaat accctcaaca atattcatgg aaatcacaca 120
aaggactaag ttcaagccta ataatcactt catgcaatat tcttttatcc attttgactt 180
caatgcttta tgggaaccca acatcattac atcaccaata gcattccaca agaaaccgca 240
ttctaagggtt tgatgtaaaa taaaaatcta ccaataccat tcaatttgc 289

<210> 30480
<211> 461
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30480

nccgggggaa ttttgnangg attncnnata ggnacnncgg nganaaaacn atgaagccaa 60
agattaattt tttctnanca aancennagg ggcgggttca gaattaggct gcatactata 120
aaagtatncc ccataccgaa taatcaaaaa taccctcata cctttgagat ttaaagcaaa 180
cctcttaaaa agtattcagg tatccggcta tgccaaatcc ttgaatgctt agaatcctta 240
ngcctataaa gcctctcaaa agtattcagc tatttgacta ggtcgaatac ctgagcactc 300
aatcattag gcttataaag cctctcaaaa aggattcaag atatgggtaa gctgaatacc 360
taaactctta gactccttag tcttataaag ccctaaaaaa tatcatggat tcgactaagc 420
cgaatagctg acactcanaa tcttaggcat tatgtcctca t 461

<210> 30481
<211> 240
<212> DNA
<213> Glycine max

<400> 30481

tgttttactt gagaataaat cacttaatca tatgagtgtc tgagctattg cctatctgct 60
ctccgacttt ggcatagaaca aaaagccgag tgcgtaagac atacatgatt tagaaaaaaa 120
tcgtccacat agcgtccatt gtgcaatcag tcataaaagc atctggacta atcatgaagc 180

aggacatgag taaaccactt taaaatataa accactactc gtatgacata actcataaat 240

<210> 30482
<211> 461
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30482

gggactaggt agacatcgaa ncnnaantt agatacnatc ctatgactng acactcagca 60
gagcaagctt tgatcatttc cttaacacng ncaaaggaga gagggagcaa taatgaaaaa 120
aacatgacac ttgggtccaa tgatgcacaa cctcacaagt acagggatca gctcaggcat 180
gacacatcaa aggaagcaca tccactcaga cacacagata agccccccaa atgaaccgcg 240
gggttactcc cactcgcatt cagaaatcac aacaagcacg aaacactaag gtcaaatagg 300
ctgaaacctg aatgggctgg ccacacatct gtgcttttct agacatataa aacactcaat 360
gatcaacgag agcaagaaaa tgcagttgac ataacgggca cttatacctc caaaccattt 420
tgtaacaagt tccctagaga tgaattgacc catcatattc g 461

<210> 30483
<211> 201
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30483

agttttgatt tccttttagn agggaannna tgcgggggcta agttggagcc aaacccagtt 60
tccctcatta agaactagct cttttcttcc tctattgcct ttaattgaat acacctttgt 120
ttggttctct atttggttct taacctctc atgcaacttc ttacaaaact ctgacctaga 180
tttcccttct ttatgtataa a 201

<210> 30484
<211> 233
<212> DNA
<213> Glycine max

<400> 30484

atgctttgct atcacttggc cacctcgtac catatatgct tacttttggg ttaacataac 60

tgacacaatg tcacttactt cactaacctg aagccaagct gaattatgga gaagggagga 120
 aaaataatca ctcaaaatgg ttcaaaaaaa caatgaccaa tatggaacat tcatgaaatg 180
 aatgctagtg aaagagatgt ctataactca acaatagaga aagtgaagat act 233

<210> 30485
 <211> 374
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30485

ttccagcctt ttatgccang gttttaatcc gaagtccaaa cacttccttg tgctttttga 60
 cctttgtgaa agtgaccttt ccagggatat tccacggagg cccttctagg ctcttctata 120
 ttggactttt cttgaattca aatggttagtt attcanacgt aatagagaca aatggaattt 180
 gaataanaca gtacatgtgc actttccttt tctgtgatac ccagtccttg agagactaga 240
 cacatgaatt tatcgtatga cagtgtgtta tatttgatg aacaagacta gatgcttact 300
 aaataaagag agctgaacac tagattaana tagagcatac tctatctagt tgtgggcgat 360
 attcctttaa cata 374

<210> 30486
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30486

tttaaaacta gtcactnaaa attttattat ttttgaaaga atcttcaaaa acaagtcact 60
 tgaaaattgt gactttggaa agtattttca aatcagcact ggtatcgatt acccttaagg 120
 tgtaatcgat aacacatcaa cagatgtgaa cttcattttg aattnttgaa aatcttaaac 180
 atttaaaaca ctgggtaatc gattacatga ttatgggaac tgattacagc tttgaaatag 240
 tttaaaaaaa tgctgggtact ggaatcgata ctactttggt atcataccaa gagaacactt 300
 ggtaaaattg ggaaacttat gtctactaat gtttgaaaaa gnttagtac ttatcttgat 360
 tgaagcttct cttgattctt gaatcttgag tcttgaatc 399

<210> 30487

<211> 347
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30487

agctaattaa tgctaaccac taccattgca catgtctctt aaaccgtggg ggaaacacat 60
 gaaattcccc accacaatct acccgacttc gacccttgcc tcggatatgc cactgaaggg 120
 caagcagttg gtggtatacc cctgcaaaac acttttgagg gccctcagt atcacccaaa 180
 actacacctc ttgcattcca caacaagtaa aaaccctcat gctatggtag aaatgggaaa 240
 gttggatcat ctagaggaaa ggctcanggc cattgaagga ggtgaagatt atgcctttgc 300
 taacctagaa gagttgttcc tagtacccaa tatcatcacc ccttcca 347

<210> 30488
 <211> 489
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30488

nnnecgttct aatactangt natctancat anaaacacaa gcccgcttg gttanacatg 60
 aatgggacct gaattgggaa cttgattata tatttcggcc aaaanggaag gagggaaaaa 120
 gtggttttca aaatctgcac tttatgccga attttgcttg tgaaatgtgc cgcagaattt 180
 tgtattagt ccaaaaaatg cttggtgatt gctggatgtg aaaaggggta tacctatggg 240
 ggtctggaca tttgcctacc gaatccaacg ggtaaaaaatg agacttatgt actagagact 300
 tccaagtaaa ttttcgagtc gatccaaccg tttacgaatt ggaacgaagg aaatgttact 360
 ggtgtatttg tatgtgaaaa gctgtgattt tgagttgtgt tttgggtaga gttttctgcc 420
 tttgccctat tttgcttgtt ttggtagtct atgatgattg gatgtgggaa tacctcgatg 480
 ttgtggaag 489

<210> 30489
 <211> 368
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30489

ggagacgacc cgaacacagc ataaaaccgg ccccgggacc cataaggggc cccagggcgc 60
 cccccctttt gccgaaaccc ccacaccncc gaaggggggg acgaaacacn agcgaaacac 120
 cccnccaca caccacacgg aagacaacaa caccacagcg cacaccnaac gaggaacaaa 180
 agcaggccac ccacaggcga gggcaaccca cacacgcaa gaccgccga ggcgccaaca 240
 gaaccaacca aggcacgacg agaaaggaaa aaaaagcaac caccacacgg aaacaaaagc 300
 caaccggga ccagcaagca aggaaccca cccgccagcc aaacagaaga acngcaacac 360
 gaaccacc 368

<210> 30490
 <211> 412
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30490

aatttttttt tttattcann cnnanaanaa caagggcccn ngngggagga agcagaaacg 60
 natttttttt ttggaccgga aacacgggag gggggggaac ggaagcacc ccctccccgg 120
 ccnngaacc cccacaatac accaggccat agaaaccccc cgggtggaaa gcaaagtgtct 180
 aaaacaaaa tagctttagt caaacgggag gaaatcgccc ctcgaaaaat gagcaactga 240
 tagaaggagt ttccttccaa tcaaagtatt tcaagcagtt gaggcctgct aacataataa 300
 cctttaactt gaacgcactg ctaggttagc gcccctcct gtacagggtg caagaggttg 360
 gctcttcct atagcatatc gatcctctat attgtttgac agtaactaag tt 412

<210> 30491
 <211> 347
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30491

gtgctattga gatgggttta aactttactt caccagagaa gcctaatga gccttgaagg 60
 ataacctagt ccaaggtcac cagatggaag cttaatagag gagcatgac gttgattgat 120
 gtaagaaaga aggctnctta tacatcatgt anggaaagat atgcaaagg gagacgaatt 180
 ntgctcaaga tgccccaaa gaattgtgac acaagagatn gngtcacatg agtatgaaag 240

gttnngggagt tctagcaaat gatcactttc canacatana ngagcagcca cttgaatcct 300
 acaaagattg tcttcangta aacgatcagg gtgttttcaa aaattgt 347

<210> 30492
 <211> 395
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30492

cccaaagcta catctccaan aaaaacncca cggggcccaa aaaagaacaa aacagntttt 60
 taccaggcgg gaaggaacac ggggggggggt cgaagaaata tccacacctc tccatatccg 120
 caaaacaaaa aaaaagaagg ctgaataaca aaccctggcc aataactaaa aacacgaaaa 180
 gaaaaccag ccgaagcaat aaaaggacac atgacacacg atcaactaaa aaaagcaaag 240
 accatacagg tcacaaaaac ctatggtcca tcaggcgtga ataccaaca aattaacca 300
 atcagagaca caacccccaa aagtccgaag tagagcaaaa gaaagacctg gccagagaag 360
 ttataggaac taatcaaaaa ttgggcagac ctacg 395

<210> 30493
 <211> 337
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30493

acctnttttg ccaatgngct actttcgcca ggaaggcatg gacaatcgca acttgtgcaa 60
 catgtaaaac acanatggga tggcttttta cagccaggaa caaacaactg aaaccagct 120
 atttcttgcg gattccgagc tgtcaacttg caaaggaaat acgccaaaac ctttagtttg 180
 atatgtatgc agtcatctat ccaaagctta gtgaaagcca tgttcaagta aactgcatg 240
 gttatgtata taatgcaact ttcttgtagc acttcgcatg ttggtatact tatattaata 300
 aaatatgttg caacagcttg gaagaaatta agtcaca 337

<210> 30494
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30494

ccggggggatt cccgagatnc cnnataanac cngcagagcg aactaggcat tgcgcccacc 60
ttacttttac ccccgcccac aaaagcgggg aattttcatt aactactccc cctcctggag 120
cacgtacacc aagcaccat tgttccacac ctaacgtcca taacgtaagg atttcggaga 180
cgaacgttct aatattactg tctcttctca cataactcatt gaagtgaatc cagtcgatga 240
tttcttcac cacgaactgc atgacgnttt ccatgcacat cccttcggac caaacaccag 300
tttgtaccct ttcgacacaa cctatacttc tttaaccgtg ttcagagcat cgcctacatt 360
gagcaacctt gggctctgaa gtaaaaaaat tcacttgcaa cggttcgtgtt tgttttact 420
cttggacact cgagaagact tcctctcgaa g 451

<210> 30495
<211> 497
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30495

ttttnttccc ggtttcgntt agcttntnt agngcgattn ncgagactcn gccngnccg 60
gggtatctct ttnagagcga cactctgcag cgctgtcgca cncnnttgn ggataangga 120
cgannacacg aacacngcgc gggaggactg tttgtcatga tatggaatac agcatctatt 180
cagcatcctg ccattctctc tatgcgcgtg attcgcagcc tgtacatggg atctctcata 240
tacaaggcat tgcgacactt tctacttggt aagcccacta tgtagctgca ctttcttgca 300
catgtcatgt cacttcgcta agacgattgt tggatgaact tcccttcccg cgcaaagaaa 360
gcacaatgat ctattttgat gatagatctg cacaacatct ttccaagaaa tcagtgttcc 420
cttaactaac taagcctatc gataccaggt atcattttca tacaccatgc cattcccaaa 480
gagatagact attgacn 497

<210> 30496
<211> 422
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30496

ntcattatcg attgttatct aacatnaana aaaccngcnc ggaacggacc taccgaaaag 60
agagcaattg aatcagacga caacnggaca aggagcgag agagcagnng aaccaccccc 120
cgcaaanggg gccngcaaac ccgcgcnagg aaggaagaag ctcccacnag caccggaaaa 180
aagacccaag acaannnaga gggaaagaaa aangaaaggc ctgagaaaag aacccccaaa 240
aaaagaaaga ggggcagcac catatacaaa gggagggaga atgggaccat aaatgcaaat 300
gaacagcaag ctcaacgcaa cgcttacaac ataacaaca cagtatatta ttttaacata 360
cgagttaacc tctggccaac aatagagatt gactaaaagg agatagaccc atagtcgaaa 420
ag 422

<210> 30497

<211> 290

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30497

agctttttatt atttttgcca gctgcatcan aatgggaaac aactggaggt ttttgagtct 60
ttgcctcttg gcattgattg catcaactca tgctcaactt cagcttggtt tttatgctaa 120
taatgcacca aaagcacagc aaattgcttt gaaatttggt catgaccata tccataatgc 180
ttcatcacta ccaactgcat taataagaat gcaactttcat gactgttttt gtaagggtacg 240
tgcttcaatc ttttaagcttc tgtcattttt acttaacaca tacaatgtta 290

<210> 30498

<211> 401

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30498

acacaaaaac cagcttatca gacaagaagt aaagatatcc aagagtttgt ttatacgtcc 60
taagcttaaa gggttattta tagaaggaat ccattgaagt acaaagttgg ccaaaaatta 120
agtaaaaagt tttttcaaga aatttactct cttgtaatcg ataccaaagg atgtaatcga 180
ttaccagtgg ccaaaactga tttacgacag ctattaacat ttgaattcaa aatttgcatt 240

gtgtaatcga ttgcacatat atggtaatcg attaccagta gtttctgaac gttntaatc 300
 aaagttttaa gcttttaate gattacacac atactgtaat cgattaccag aggagttttt 360
 cagaaaacat tctcaacagt ctcattttt tatctgtttc t 401

<210> 30499
 <211> 76
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30499

agcttttagcg tgaaactnta acttttcata ttctttcaat tagatatttt taatattggc 60
 cttttattta tctttt 76

<210> 30500
 <211> 387
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30500

tatagtttaa atcaaaatag atctttgtat aatgggtgta ttttgtgctc tttntatgaa 60
 tcttcaagag ttgtgggttaa ggatttcccc ctttcttcta agtcttgana atccttaaga 120
 atanttttcg tccactagat acctttttgt ttagtaatgc cttgacctcc tcacaaacaa 180
 atgtgtcttg tacatcacta gttgaagtgg tttccaagaa aaaatcaact aaatgagctt 240
 gatgagtttt ottaagatcc ctatgctctc tttcaagatg ttgaaaactt gtttagtttt 300
 ctataagctt tagagacgtg agcatataca gataaaagtt gattgtaccc ttttaattaaa 360
 tattttctag attcanatat tatctct 387

<210> 30501
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30501

nnnaagtat ctttgtatcg taccnggntn ntagannann cnncccggag gcacnaaaaa 60
 ggaccctcaa gcagcgcgca cncgtttttt tatgcgccaa aggaagccca ccaggggggg 120

agatattcgt tatctaaact ctcaaaagtg actgagaccg tgatgaatat agacgagttc 180
gagtcacaat ccgaatatta ctctcgaata tcggccgtag ccaggcggca tgaaagaaag 240
gcgcgcaatc ttgagaaggg atagaacggc aaaaattctt cgactttgct ctccaagtct 300
cactgaatgg tgcccccta gattgaaacg cagatgttac caacctgact attgatcaac 360
tcatggcgac ccttccatcc tattgaaaca ctccaatgga cacaacctag acaaagccg 420
cctcagagca cgacctcggc tccacgaaaa gaaccn 456

<210> 30502
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30502

ataagcagta cagaagcann cnatagaaac nccgcgagnn cacacgcaag caagaccaat 60
ttattttttg ccgnaacaa cagccgaggg gaaacacaat aaacaaccga ctggacacaa 120
agaaacaaaa aaaagacaca cgngtgtacc cctatctgca cagaacaaca acatctaaca 180
tagctttact cggaggaaca cgaaacccca agaggacaga tggaactac cctatgatca 240
acctggagaa tatcaatggt gaggagagag agactatggt ggaccagaat caacatgctc 300
gtcagaaccg gtgagttttg gagggaaacc acaaccacaa actgctttac aaaacctttg 360
gcatactcac tcaaagttat ttagagctag gcatccgcac cgtaactgtg cctaaagggt 420
gacatataaa ngaagacaac gaacggggcg cn 452

<210> 30503
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30503

aggagtgnnn ctgaacaccc agtaaccgac cccgggatcc caagacacca gaggcggcaa 60
atTTTTTact ggagcacaag gaaacagccg ggagngaaga aggggaccaa cacaccang 120
ggcccaacag gaacaacaac ggcgcgagc cccagaagca ggaaagagcc ggccacaccn 180
tccacatgta aagccagaca acttcctcct tagcccatga cccacgacat cgcgtgcatc 240

cacagatcgc atcccaaata tccgaggcac aacccttttag accgtaatat gtggcggaac 300
 caggcgagca agctgagggc gggcccacaa agcgggacct cggactggaa caatagaaag 360
 gcggtatggc acaccactag cctctatcga cagccactca tgacacatgt gcgccgcct 420
 gcgcaccttg cgacg 435

<210> 30504
 <211> 480
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30504

agggatantc tgaatcttga nncnnnnnnn nnnnccgagg nngcnnccagg gntgggggag 60
 aaanctttna tacaattccc accngnggan acccaccagg agggntttca tatgaagcat 120
 acatcactgg ggggtataat tccaccttat tgaaaaattt aatctgcata atatcacttg 180
 tgctcaattt cttgtgacca agtaacattg cacaatccat ttcattccaac atgtcatggt 240
 tgtggccaaa gcaacatatt gtacatacca atgatcgta cttaaagttaa cgtgaccatg 300
 aaacctagca ttgcaaccac cttgtatcat gttttcctca cgcttcctag ttgtcaatgt 360
 cagaccacta tcattctaat atccaacatg agagcanata aaaaagttgc tgtaatgttn 420
 tccctattat gaattctata acagtacttt tacggacaaa ggtaccatta tatctagctg 480

<210> 30505
 <211> 141
 <212> DNA
 <213> Glycine max
 <400> 30505

agcttttctg ttaattcatc tctgttaatg gaaacgatgc ttattaaggg agtagttgaa 60
 aacaccctgt atatcactag accctgtgaa tgaagtgaat tttatgcatg ttaaagtctt 120
 cttatttttt tttgaaaaac a 141

<210> 30506
 <211> 349
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30506

tcttttnggac cttgaacaag caattaactc ctctttttaga ttcattgctat gngctcgcga 60
ctgggtctctg tcttcccttc gcaacttgag ttcactattg ctaccccata gagctccgcg 120
aaatatgttc cggccatact cttccttgtg agccctcttg gtctcttggt caagggctct 180
tgcagtaatt gcattctctt cccgtaaccc ggcacactcc ttcggaacgt gtgtagcggc 240
caacttgaac ttctccttgg caagtattgc ctttccctaac tctcttttga gagtttggac 300
ttcttcgtcc tcttccggtg cttcaaaaact ctcttcgctg acgactttt 349

<210> 30507
<211> 157
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30507

atctatcata ctttctctcc atattanctg agtccttcat aaaaatattg gagaataagc 60
tgttctgaaa tctgatggtg ggggcaactg gcacatagtt tcttaaattct cttccagtac 120
tcattcaggc tctctccact gagtngtcta atacctg 157

<210> 30508
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30508

tctttgtggg ttgatgaact ctatcgcaca tattgtcttg atcattgctg acatattctc 60
aattaggtca gttgcctctt caggaatctt taacttttat ttttccccct gctgaagcat 120
ctagcagttg ctttggtttg tgggtctcagt cctctataa acatattcaa ttgagttggc 180
tcagagaacc catgggtggg agtctttctc aataaacctc tatacctctc caacgcttca 240
ctcaaggact cgtcanggaa ctgatgaaat gaagagatag cagctntccc ttctgtagtc 300
tttgactttg ggaaatattt cttcagaaac ttttcaacaa cctcttcccta aggtttcaga 360
ctgttacctt taaatgagtg 380

<210> 30509
 <211> 300
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30509

attccaacct tttttggtgg ttaagttctg gatattatcc tttccacca ttctgaaaat 60
 cattccattt ctgataatg ccctaaaata gaaattcaca tcagaaaaac tgtgaccatg 120
 atggtaaacy tcaacagaga tcatcacctt ctagattctg tatgtatata ctctttttgg 180
 ctcgatataa caaaaaatac taattaagtt ggtagaatat taattatcca attcatttaa 240
 agataaccct tttcataatt atcttcttgn acttgattca gaataaagat gcttggaaca 300

<210> 30510
 <211> 481
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30510

nnaaagacaa gcgttcaagn agnatcttca caccatanan tactcaagcc tntanaaagc 60
 atgcaaatgg aggaaggaga ggagcatctg gatatatgtt gtttctactc atgaggtatt 120
 ctagattaaa gaataccttg atcctcaact agttgccaag aaaattctca taaacctaca 180
 tgatattata ttttttggtt tatgccctac attaaaaata gcccttagtg cattgttagt 240
 ttggggggtt gactgtaaatt tattttggtg tccttttgtt gagttgactg aacgaaaatg 300
 gagaagagca ctacacagag gagcangatg aacagaatgg agaggaatga agccaagatg 360
 aggcagttgt agtggatcag attctacgat ggtgcatact cgtaatggta caggctgaca 420
 aagtgggtcg aatatgaaga gcccgtcagc taagcattgg tgggggggtga aacaaaattg 480
 t 481

<210> 30511
 <211> 325
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30511

cggggaccct tatccannaa nnaaccgctc accctccatn agtcntttac caatggacca 60
 tatgtcaaac aaacggctct tgtgaataat tgttnggacc cggtgtacag ggagggcttg 120
 ttgcaggaca agattattgc tatctatgtc aacattaaca atcaatgatt ttttggacaa 180
 atatcaagcg gactgaaaac aaaacagggg taatgaataa aagaggactt tatcgatcct 240
 cttttaatct tcataattgt ggcactatgc attggacca aataaaaatc ttaattttag 300
 tatggcctac caaaatagta taccg 325

<210> 30512
 <211> 180
 <212> DNA
 <213> Glycine max

<400> 30512

atatgcacat agaggtcata cagcaatcat agaaccaaac ttagaccaag catcatacaa 60
 atgagtaaaa atatacctcg aatgctgtca caataatcac agccagaaag aatgcataag 120
 tcaataaaat tgggtccatgg tcatatttag ctccctccaaa atctgaaacc aaataaatac 180

<210> 30513
 <211> 237
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30513

tggtttccaa tttttctgat gccacaggtc gggatatgga taagattata aaaggaaggg 60
 ttgaaaaatc tgaaaagggg ggcaaaatgg gttgctttaa atttgtagt gcctccagct 120
 gcaacatgct gttaatgtgc cattgcactg ttataacata tgcataata ctacaagtaa 180
 aagtcttgct ttatttaaac cttttngtgg taaatctgct tattagaact caatatg 237

<210> 30514
 <211> 492
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30514

gcgcgatatt ctgtanttca acacaanaaa cnaagcnggg aggaactnga ggcgaaaaac 60

ccagggaccg tcattttttt acacggaacc acaccaggng gggaatggga tacaatattt 120
 tacatcgaga cggacacatt tggggatact aggcgattca ggtatactaa atattaaaaa 180
 tcgaattagt tgaactttta atttcttaat taaatccttg tggaagagac gaataggcct 240
 ttctttggag tatgataatc actgaacctg tgagtcacta tctttataag attcgacagt 300
 cacatactca atatgatata ttctcatact attaagtgat tatatttatg cctctagtga 360
 tgatgtggta atgagcatat atgcatcatg cagattacat gcatgcacgc gtgtaaataa 420
 taatggaacg tgccatgtgt catgtgcttg cttctgtcga gttctgaaat cagacattat 480
 ataaagttgc tn 492

<210> 30515
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30515

aggggacctg actcgcgatc gagccgacgg ggatctcaag gcaccgaagc agcacngtgc 60
 caaagcccnc acgcnccccg agaggggatcg gacaaaaaaa acccgccacg aaacgaccta 120
 ggctataaaa gaaagtacaa caaacacggg aggcaacgac aacgacacac aaaaatgaca 180
 aacgcaccgg ggagggggaac aaaccgacgg ggaatccagc taagaagcag cgggacgccc 240
 atcatccagc aaccaagctc tgcaccatag aagcaaactg accaacggaa aaacgcaggc 300
 accacctggg gggaaaacaa cgaccgacaa ggctgaaccg caaaacaagc cagaacaaca 360
 acacaacca ggccggccac agaagaggcg gaagaaacgc gcccatacag gcc 413

<210> 30516
 <211> 303
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30516

atatgaacgg tataagatat tataccaaat aatttttgtt tggttaggaa ctttataaaa 60
 tagttaaaaa aaaaggaaac ctgattataa aaaaaagata tgatttttgt ataggacct 120
 aatagcatat gatttttcta tagaaacata ataggatagg ataaaatatg atntttattt 180

gctctagaat acagaaacgg tatgaaagaa aaaagaaatc taataggaat agaaaaagga 240
 ttaacatang anaactaata aaaataatga gaaatataaa ggaacaccgg actcacatct 300
 tgt 303

<210> 30517
 <211> 223
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30517

tatggggacc catcacatgt ggcctaggtg gcggtcggca atggtcacaa caagtgttcc 60
 acatccacaa tgcgcgcata aacccaccat cccctgttgc ccacgtcaac tgagctcacg 120
 tactcccacg tagcccatat ncctcgttct ctcaacaccg ggtccccatc aatccttcca 180
 agcttncaca acatccaagc caaacaacat tcacacagca caa 223

<210> 30518
 <211> 460
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30518

gggtgctgan ccatgagact actgcaencc nattngnnaa cnaccctgtg tggttgcgcc 60
 cggcactaca cgccgtgccc tttgtttaat ttgaatccaa gccccatcct ttcggggggc 120
 aatgcccctc tcattacctc tatcccgggc aagacgatga ggatggagat acccatcttg 180
 gccgcctgct ccacctcaaa gatccgtccc ccatgaacta cccaacgaa catagtccgg 240
 cctatcccgg cctcacgcta acccgataaa gaatatgatc ctttcgctga agatagggag 300
 agatcgaggc gcttgagaga ggttaaacca gtcggggcct tgcaataccg atatatgcct 360
 actcattacg atggcgccac aatcaattca tccagttcaa ggccggcttg tattacaaag 420
 gacactgtcg atgggctctt cgatgtttgc gaagatgggg 460

<210> 30519
 <211> 349
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30519

naagcttttg tttgggtccta tggcatngnn gnnngcatcc ttggatacat ttcttagtgg 60
ataatnttgg tcctatagaa gaaataataa aacaattcgg agatattctt gcattgtata 120
tccccctgc tgctgtgtac agagatccat attctattca ggtgtttgaa ttttagaccc 180
tctataaata agatttttca tcatatcagt tntctaagt actaaatata gagaagaaat 240
tgctctgttt ttgccattta atttatagaa aacatttaaat cttctcaaag gggagtaatt 300
ggaaagaatg ggttgcttaa tttctacaaa ggctgataac tttttttct 349

<210> 30520
<211> 307
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30520

catcaacagc actgacccaa ttattttgta tagaagggtc ctctccttc cgcgagacca 60
cagcccttgc gtactcaatt tccaatatcc tttctataa ttgattattc aaaactgtta 120
gctataatgc tatgacatca taacacgtag taacaatggt ttgtttccta tgtaagaaa 180
taaagattag agaaattgat aagttattgg ttgtataccc aatttcaatc cctggaattt 240
caacaacttc cttgtaagt actaaatggc ataaggttgt agatcattcn gtttcacatg 300
attgaaa 307

<210> 30521
<211> 320
<212> DNA
<213> Glycine max

<400> 30521

agcttattat atgatgcagc atatttaaca gttttcatag gtgatataag agcatttcac 60
aacgaacatc atcttttttc tctagattct caaactcata attctcattt tgaaatttgt 120
tgagtgttc cagaacttca tccatagaaa gacttaattc attgtctcct tgcacacacc 180
gaaaggccaa ccctgtact gaagttagta tccttttaac tacttgggtc gactcaaacc 240
caaaggatgg gtctacaagc tctaagct ttactttttg cacctttttc atggcaagat 300

ttgccaagta gcttcatctc

320

<210> 30522
<211> 428
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30522

nnccccggac taacgtccgt cntacggaca ctatagatac tcagcttgag ctggagaaaa 60
ctggaggagg tttgggtttt acttgctttc tccctgagtg acattgtatt ggtggatatat 120
gagtgttcat ctagaatttt tgtgcatctg catcatatga atagtgagac aaaattttct 180
aagtagaaaa gttctcagaa gcgaaactct ctatttaatt gattacaccc tatcgtgatt 240
gttacccaag tgtctgagct tgccggagtat gtctataccg tttaatcgat atagcctctc 300
gaatcgatac aaattgtgat gaacaatgct gactattcaa gagttctctt tatcgatacc 360
atggaatatt gacactctct tcatagcagt caaactccag tgtctatctc ataccttgca 420
ctacatcc 428

<210> 30523
<211> 276
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30523

agctttaacc tttggccatc atttctgccc caaggcgtga aaggagagca ttttcggcgt 60
cgtgaagtgc gtggctacga gtgggacttc gaatattcag gtttgggtgg acttctttct 120
ctcttaaatt tcgtgggtat ggggttntgg gagatatgat gggtagtctt gctagggtttc 180
tgctgtatga tgattatttg tgaagaaatt tgttgaaagc ttggtgaaat cgccatgttg 240
gatgagttaa acatacccat ttctgtttta ggtttt 276

<210> 30524
<211> 400
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30524

tgcgtagccc accatctttt catagnagag ttattttattt gtgtctacca tcacgatcat 60
 cggctccctt tccatcattg ngggtaccac ctgngccgcc agatccctcc accttttggg 120
 cgtgttcttt gaaagatecg tccccctttt tgcaaatgtt ctgtaattgc atcctatccg 180
 gaaccatata aaaattgtac tgatactgcc taacaaaggc aaccattang tccttccaag 240
 aatggactcg ggaagattcc aagttagtgt accaggtaac agctacccca gtaagacttt 300
 cttggaagga atgtattagc aattcctcat cttttgcgta ttcccccatc ttctgacaat 360
 acatcttttag atggttcttg ggacaagtag tccccttgta 400

<210> 30525
 <211> 321
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30525

agcttttatn taaactccnc aaggaagnen ctcgagagag cttatcaaag aagcttctca 60
 acgtaactac ctatgctata aatagaagca tgtgtcacac ttgtggtaac tctgatgaat 120
 gagagtcttg tgagacacac ttcaaagatc aacttctctc cttcttttcc tccttcaatt 180
 tcctgctccc cctctctctc ctctctcttt cttttcctcc atagaagcat cctctccaag 240
 cttcttatcc aagcaccttc ttggtggcaa atctccttct tccatggcgt attccttagt 300
 catatgccat gacaattaac a 321

<210> 30526
 <211> 227
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30526

cacactgtat taacatgaag ctgacgattt caccaatccg gatgaaagta tttgtgagtt 60
 tggacttgag tgtttgtgag ccaccttgat gtcaccctaa catcaagtgt tggacctgag 120
 tgtgtagaag tgatctctat tgntcagaga gcaatctctg gtgtgtattt gatttaattg 180
 tatacaccgg agagtgattg agagggagtg agaggggttc tcatatc 227

<210> 30527
 <211> 300
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30527

ttcnttttgg nttaagctca cattggcatc acccgcataa tatacggact tcaggtaagg 60
 cagtctgcaa ccacgttgca taactgacga cctaaataag ttgtcatgtc aaacaanatt 120
 taaaactaaa atataaataa natttaacta atttcgtcct attcaataat atcatatgta 180
 aaaaacctta ttcaccaaac aaaatatcta aaatatacctt aatttaataa aagttttattt 240
 taactacgtt ttttcctgtg cgagtaatga aatgatactt aaaaaatatt aatttttaatt 300

<210> 30528
 <211> 477
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30528

nccttagggg aaattgccga ngtatnnctg ngaanaatan annacncaac gctaagggca 60
 cagggtacatc gtaaggaanc cgtttgattc cttaccaaca gnngcgacac ggagggggcgg 120
 cataacatca taaacacaca tatctcagat accttaattg tgcaatcaca atttacacac 180
 gcatgacgat gcaggggacta ggtactatca tgcccacgac ggcgtatcga gggcgcccac 240
 ttcttgacta cagaggaaca catcttcggg ttagaatcgt ggacgaacaa tgcaagaact 300
 acaacgtggc tcaaggacaa gaaataaaaa gacttccctt cgttggggatg ggagacacaa 360
 tgcactttct tcataataga agctcactgc atattgaata cgtgcggaac aatcttatca 420
 ncggagactg taaccttaata aggtcaagag tctacaaatc atagatgtaa cctgggtc 477

<210> 30529
 <211> 254
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30529

agcttggtct tggtttanac atgattgata catgatttgg gacttgtag attcaatttg 60

ngcaaaattg gatgagggaa agtgtgatat cgaaaatctg cacttatgca aaattttgct 120
 gtcaactaag tgcagcagaa tttggctctg tgcaaaaaat gatatgaaat tgctggttgt 180
 ggaaagagta gcaccgattg ggttctggac gttttctatc agatcccaac ggtcaaaatg 240
 tagatttatg tact 254

<210> 30530
 <211> 298
 <212> DNA
 <213> Glycine max

<400> 30530

tccaaataaa atatttggga acgtctatta gactataact ctttaacagc tactgtgtat 60
 ttaacgggtt cattaataag tcagattaaa aaataatttg tcaaattata attttggatg 120
 acgcttacgt gacaaagtgg acgtttgata ttgaaaaaat taaaattact tattttaaac 180
 aatataagaa ctaaacgtct tcatttagag ataaaagact caaaatctca gattttaaatt 240
 aatgggtgaac caaaattatt aataaaaata tataatcggtt ttaaattatt tatttttcg 298

<210> 30531
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30531

atctttgccc aataacactc ctaacatcaa tagtaatatt aatattctaa ctcccctccc 60
 ctaaaccaaa gttcacttag catcacgctt gttacaaagt attagaatgt cgcaatatct 120
 ttaatgtatc tcagattatt gttaggatca tattgcaagg tgtgagaagt gagtatcaca 180
 atgaaagttt ggcattctaa tgtaaggttt attaagcctt aaccttgagt tctcaactac 240
 aatggctngc ttttgtggtg tagttcttcc cagagtctta ataattggta ttagagcttc 300
 ttaccatgtc tctatgct 318

<210> 30532
 <211> 368
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 30532

tagctatnat aggggtgtgta tagctaagct ctatcttctc tttctaagga ggtgagctta 60
gctatgagag aggtatgtgt agctaagctc tagcttcttt aggaatcttc ttaaggaagt 120
ttctcacgga ggtgagctta gttatgagag ggggtgtgtg agctaagctc tagcttctca 180
aggaagtttt ctcaaagaag cttctcaagg aagttttctc aagaaagctt ctcaaggaag 240
ctacctagtc tatanataga agcatgtgta acacttggtg taactctgat gaatgagagt 300
cttgtgagac acaacacaaa gttcaacttc tctctctttt cttcttcaat ttgtgctccc 360
cctctctc 368

<210> 30533

<211> 277

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30533

ttttgggggt ggccatttct tggatggcct tgattttctc aaggtccact tggaccccat 60
ttctaccaac tacaaaacct aagaagacta tattatctac acaaaaggta cacttctcta 120
tatttgcata gaggggtgttt ttctaagga ctgaaagaac ttgcctgaga tgtcctaagt 180
gatcatctag gctectactg tacactanaa tatcatcaaa ataaacaact acaaactctac 240
ctatgaaatc cattaagaca tgatgcataa gcctcat 277

<210> 30534

<211> 293

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30534

ctagagctnt agctccacat acctctataa tagctaagct cacctccctt gagatgagaa 60
gctagagctt agctacacac cccctatgat agctaagctc acccccatga caaaaaacat 120
gaaaataaca caaaaaagtc cttattacaa agacaactca acatggcccg aaatacaagg 180
ctaaaaccct atactactag aatggccaan atacaaggcc tagacgaagg aataacctat 240
tctaatactt acaaagataa gcgggctcat acttagccca tgggctcgaa atc 293

<210> 30535
 <211> 367
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30535

 ccaacctggt aggcctagaa cctcctccac caaggaaggc ctaacctact ggaaaacatg 60
 gccactggac cggagaagga aaaagaataa tggaaaacgt cccttcaagg aaaagatgag 120
 tcaagaggaa gctcaccacc atangaagac atgcgataag atcttggatg tatgagaaag 180
 ataattggca agagaaggag agaaagggtg cgatatcttg tgcctcaaag gaggtctgaa 240
 ctttgaaagc gaattcttaa atgatcaaag gtgacaaaat gcacacctat ggcttctatt 300
 ataccctaag ggcacaaatt tggaggaaat tgaatctcta taacaattca cttgaattga 360
 catgaac 367

<210> 30536
 <211> 474
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30536

 naacttaaga tcagatcaac ctnnaaaaag ccccagctga aacanngtca ggtgcggggc 60
 caaccttaat tatgggttcc aacaagnnnc ccaaatccag gagtaccatg aagggataga 120
 gaagacatac tgttgataaa caccacaaa atatagttat ggcaaagcat tatttttcca 180
 acctttttaa aaatataaaa ttattaataa tatttccact aattaccata ataatatatt 240
 aatggtagaa atacttaatt cttttaagt gataacatan agcctcttaa naaattgtga 300
 gcaagccaac ttgttatcta gtccgttctc atacacattc ctaggagnat gtatcatatt 360
 cattttcaaa tcanaataag aaataacata tcatataagt tcttaggtgt caggtcaatc 420
 ctattgatat caataaaaat aatttaatac tgctattggg tgtctagcac tcag 474

<210> 30537
 <211> 449
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30537

agggagcgct ttgcattgan annchnccnn nnttngnaaa nccgccngga ngganacana 60
gaaaggcaca agcaggcaac aatttttttc gaaaaaccac cccaccaaga gggggcatcg 120
gaaaccaacg cacacacccc accaaagaaa cccaccccca caaaggngga ccgggaggga 180
gngaagaaca ccaaccgcg gnnccggggag cagaangacc accacagagc acgagaagng 240
agacnctacn gcggcccca caccgcgag aagaccccg gaacaacgcg ggaaacaagc 300
aagngcccgg aaccaaccgc ccaaagcgg agaccataac cccgccggcc gaaccggaga 360
cggcaaacac acagccaacg gaaaggagg accaaanaca cccgcaccaa gggaaggacg 420
cgcaccggac accgaccca aaaaagacg 449

<210> 30538
<211> 491
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30538

gttcgtagnc atgagattca aactnnaaa aaagcnngaa anaaancaa nncnagaggc 60
ggagaactct ttttaactttt ggagccgaaa ccggggagggt gcaaatacaa attgcaaatt 120
gtctctaaat caatttagcc actagtaa atcgattacat ctctggtaaa tcgattacaa 180
gcggattctc tcatagttaa cgcttccaga atttcttcaa ctaaaaccaa agtataggga 240
tctgagggct acaatantgg atttggtcat ccgtgtacac ctaactgaag tgggtgaaga 300
atatagcccc tctaaggtaa ccccaactat gttgcattgg acttatgctc ttcttgtgcg 360
tgtttataga agaaaaactt atttatattg tacccttgcc aagtgataac atttctttaa 420
accatcctg ctactttctt aatgttcaga agcttatcct cagcagagat tgatctttaa 480
ccatctgaac g 491

<210> 30539
<211> 138
<212> DNA
<213> Glycine max

<400> 30539

agctcctttc ctttttccac tcaggtgtcc aagtgtggga tggcataggg tggaatggtg 60
gacagcctca gtaactgcta ccatatcctc atctggcatc atgtaaagag atcctcgctt 120
ctttccacga gccacaat 138

<210> 30540
<211> 240
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30540

aagattaact tcattcatta tatgatgaat tttaggggac gtttaatatc catcctcatt 60
ggagcacaag ttgcacataa acaatggctg agaacatgag acatanatgt tannggtttt 120
agaaggaaat cctgtcatca nttccaagtt attgatattt taagttcttt attttanaat 180
tactctattt tataaatttt aagctaatta gaattgtata tatatgtgga caatacatte 240

<210> 30541
<211> 188
<212> DNA
<213> Glycine max

<400> 30541

tgtgaaactc tgtctgtcac atcattactt aatggacaaa tactaatggc aggtaccaa 60
gccaaaaaat gagaaacgcc caatagagca ataaaataaa aaattaatat gtagcaatct 120
gaataagggg tatatcttag gtaagaaaaa gatattttaa ccaattattt tatgaatate 180
atcttaca 188

<210> 30542
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30542

gccttaaaaa ggggtgnttt gcaccttctc gctattccta tattctggct tagcgagcgt 60
ccgctaagcg caccactcat gggctaagcg cgaggaaaac actagaaaaa gatgagttgg 120
acaggttctt tagcgcaccg cttatctact agtgcccact tcgtcatctg taacgagaaa 180

gctgcgtaaa gctgaatcga tttagaagaa gttgactaag atcagagctt tgctgttaga 240
 ttttaaagag acaagtcaag ttcaagagtt tgaagatttt gtgctgaaat tgcggaccaa 300
 ctgaacagag ccgttgngct gaatgattgg aggatggaat cctaaggagg tcatctacac 360
 tgattntgat ctcatt 376

<210> 30543
 <211> 110
 <212> DNA
 <213> Glycine max

<400> 30543

tgtctgccgg gcagacaacc acaaatcatg tttacaaatc attatgaatt atggcattcc 60
 tcgaaagtag ccgtatgatg cgtaccacca agcgtcatag gcactatacc 110

<210> 30544
 <211> 186
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30544

agcnttaacc aagangggat ggtccatttc aagtacttga aaagataaat gacaatgcga 60
 acaagattgg attgcctagt gagtataatg tgagtactac atctaattgtg tttgacttaa 120
 ctctttttga tgtagatgga gaagccgatt tgagaacaaa tccttttgaa gagggagaga 180
 gtgata 186

<210> 30545
 <211> 437
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30545

nggcgattca gttgcaagtt ctacgnccac tatnaatact cagcctccat cattgagtta 60
 agtccaccag gaggaatctg ttaacttggc ttgatcaaga ttaggctaaa ctatcatgag 120
 gcaatcgggg ttaatatctt aggaaacaca ttaggacccc ttgaccttgg ttgaatgaaa 180
 atatttttta acttcaggcc cctataagga agccacgtgt ttctcacatc attctctccg 240

tgattttctt ttgcacagat agttacacac ttgtcatatc atgctatctt acacaccgac 300
 cctattgctg aatagcttac caatacacia gtcctcagag ttcatactcg tcttaccgtt 360
 atctactttg cgaccgggca cttgcgagca acatagccat cagctgggta ataagaaaat 420
 gattttacaa attttgt 437

<210> 30546
 <211> 193
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30546

agggcgcgca agctttcatt ttcaatcngg agncgcncga ngggggcatg actcaatcgg 60
 acatcctttg ataatgttat tgtcgtttga atttgctacg agctatcgtg gtaatttaga 120
 gcatctagat atatttcggg acacaaacag acatcctggg ataaagacat tgtcgtttca 180
 atttggtcag agc 193

<210> 30547
 <211> 539
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30547

attgttgttt ttttctttga tctacnnga cacnananan acncaagcgn ctagnctcaa 60
 tngggagagt ctcgaggatg accgtttata antctgcatc cgagnaaaaa ganagggggg 120
 cgngaataga cnaacagacc ccatcctcaa cngggagcgn cncgaaaaaa aacgggacgc 180
 aaccggacga cccgggaana aaggggaccg gccccgcaan aggcgcacga gacacgcagg 240
 cggacacaaa caggaacagc ncgcggacag ggaagaccgg aacacggaac agaccacccg 300
 agacacaacg gcaaaggcgg aaggaaaaag gagacgaagg caccagccg gcaaagagcg 360
 gagcagaccc ggacangaaa acaggaccgc agaccggaca cgcagcgca aaaaggacag 420
 ggccaagcaa aganngcgca ggagcaggag cagcgcgaac ccacgccgac gagggccnag 480
 acaaganggc aaggacgcaa gaccacccg ccgaggacaa cagcgaacgc caccgcgcg 539

<210> 30548

<211> 302
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30548

 agctttgttc aaaaatcaaa agannnaact ctgcgagaag gttttcaagt tttgtaaaag 60
 ttataactct ttccaatggt ttccattgac tagacatgaa gagtctataa aagcaagacc 120
 ttgacttgca tttccataac tttttgactt aactttttta caattcttta taacaacttt 180
 tgagaaacct ttgctaactt attattcttc ttcttctttc tttgcaaaaa gctttcttaa 240
 agtatttggg tttccaaacc ttgaaaacaa aaatgtggta ttcacttttt tctttctctt 300
 ct 302

<210> 30549
 <211> 345
 <212> DNA
 <213> Glycine max

 <400> 30549

 cagcatcatc tatggcgtca gttatcttgt caacattggt gaggttggga tcaaagtgca 60
 cttttgtcta ctccaaagca agaacaacaa ttgctttttt cacgcctact accaatttga 120
 agggcattcc caacagagtc agaacagctt gtgcatgcca ttcctttaat cctcactcgg 180
 cacactggta tgtcctgctt atgaagctca tttactogaa acccactctg ctctatgctg 240
 tcttttatgt gttgcaccta caaaaagaat atcaccaata ctttggtcag tgtaaattatt 300
 gaagtacacc aaagacttat atctattagt taatatacat aattg 345

<210> 30550
 <211> 290
 <212> DNA
 <213> Glycine max

 <400> 30550

 gaagcaccgg acgacgcata cgaccggggg acaagacaca gaggactttt tcccacaggg 60
 cgccgggggg gcaaacaacg ccaccgggag ccaccaacg acggagccag aggaccgacg 120
 agacaggcga gaccaccaca gacggacccc ccgaaggaca aaaaggggga cagaaccacc 180
 gaagggcgca aacgaacca aaaaaacaga ggcaaagcca aagccccgaa gaaggaacaa 240

gaaaacagca acggaaagga gagacccac acggccagaa aagcgcccc

290

<210> 30551
<211> 303
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30551

ggtagncct gnangatcaa atgnaaacgg ggaaagagga agaaattctc tttccccgga 60
aaagcggggg tattaaactc tctctccga tctaactgta ggaataacat ggcgcccgtt 120
cctcacagtt gattatctag gagaaaaaat ttgctttctt aaactagcta aatatttcag 180
attccgaaga acatgcacat atctactaca tggttaacta attttattgt tatccgtaa 240
tccacaaatg gaactctgcg tcgaagcttt ttggtgctat cagctcaata ggacataatc 300
tca 303

<210> 30552
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30552

agggacgtct ctttgtanan caccggcaat ncagctcggc cccgggatcc tctagagcga 60
cctgcggcat gcaagtcgtc catcataccc gaccncgca ggggcgagga tgatganaca 120
catcttccga ttcaattgtt acatcagata acaatctatt tatatttaaa atcttgggtg 180
tatatagtcc agaactttgc taacttccga tagcccgag tattagcata gattgagatt 240
aagaattcgg agggatacgc tactgatcgt ttgatgatt gcactgaata tgtcgggcta 300
cttggcctga cgcataatagg gagccaaaaa agccgcgata ctctgcggca taagaacct 360
tctctttatt ggctccatt ccaccgaaca catggggaga tttccttcca tgtaaatact 420
g 421

<210> 30553
<211> 484
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 30553

gcacnagng aaagtcccg nagatnntct nnnnntnaga nacnnaacnn cagagngaaa 60
 caagatggaa gtggaaggac attaccttat ttgtccaacc acggacaaaa ccgagagggg 120
 ggcgaactng gaancaacct atcaaacagc nccnganncc accaaacggt gttggatata 180
 catactttat ctcaccctcg actccattta ggatggctga attattcagc gaacttgaat 240
 attattcttc aacttcaccc tctgcataca tagccccac ccttattcat gatgatgctc 300
 acgactaaaa ccaagactac tatgtgccgt tatatgttaa tagagaacac aacgaccagg 360
 acagttcctt cgtcgaatat ctctgttata tacgaataat gcgaccctga gcacaacccc 420
 gaacttatat taggtctctc tgctgggtac ataaagagca aaaatgaata aacatattct 480
 tatt 484

<210> 30554
 <211> 447
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30554

aggagctatt cgtttgacac canccnnnt ggaaacacgc gccgagatcc ttagaggcac 60
 cagcaagcng cagctttttt tcatggcctc cnanggcgga gagcggggtc tatactcatc 120
 ttcagctcga agcggcgtct cctctgtctc tttcttctcc attctgcagc cgttcatctc 180
 ccagaagcaa aggaatccat tgatgaaaaa gatcctaggc ctacaagctc caatggagct 240
 tacatcatgt ggcacaaaga tcatttttga ctacgtgatg ttcatttgcc tcctccatcc 300
 tttgttccg tgcattctct ataacaacgt gagcttcac tttatttcca tgtatatcct 360
 ccattgtcct gtgggcaagg agaagggtac aaaagactcc acaaagataa atcgattata 420
 tctaaatcta cacttgtcta gcattac 447

<210> 30555
 <211> 518
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 30555

naatcttacg tcangngtan gtacnantnc nananttnag annacnntcc ngcgccccaga 60
gaaggagngc acggaggaaa ngcttanctt cnctanaata cngganagca annnncgaac 120
gacgggcgct gttggccaac acannaaagc accacacgaa gggcagcncc acccaagaag 180
gccnaacctc gccttganac gaaggacca ganngccct nncacctacg aaanancaac 240
tttttggtgg aagtgtgtga gggaacaacc tccccactga gtgtgatcca cgaggcgctc 300
caaccagaca tactgtaggg gggggttaat atccatttat ttggaagggt aacttgacag 360
ggtgtgaggg tctatctgta ccgggagatc gatcttcccc cttacctttt tgnngggacc 420
gtgcaaggca cgacccacca ttgacttcgg cttatgtggg aacattgaat ggaaattctc 480
caagtgtctt tagcttacgt taccagaac catatcan 518

<210> 30556

<211> 332

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30556

catgattggt acatgatnta ggacttgtat gattcaatnt gngcaaaatt ggatgaagga 60
aagagtgggt ntcgaaatct gcactttatg cagaattttg ctggttgaat gtgcagcaga 120
attttgtata agtgcagaaa aatgcttggt tatggatggt tgtgaaaagg gtagtacata 180
tggggttctg gacattttct aacagatccc aacggtcaaa atgtagactt atgtactaga 240
gacttccagt aaaatttttg agtcgatcca acggttaacg aactggaaca aagagaatgt 300
tactggggta attgaatgtg anaagctgtg at 332

<210> 30557

<211> 353

<212> DNA

<213> Glycine max

<400> 30557

agcttgtcta ttataatta tattgagaac aactgaggag tgttgtgttt tgtacaattc 60
atacataaag tatgtgttaa tagacttctt ggattgtgcc tgaatgaaga ggaaaatgcc 120
ctgaccgact cttcagagtc tacgtcttgg ggataaatac acccggtttg agtacttctt 180

tatgcttgaa ccaatcccac atgattggag cattctactc aaacaacgtg accctaacta 240
 gtctccctat gattttacct agtgagtgc ctaccctacc actgtgtgga ttgttatggc 300
 atgcactcct tggcaccgga cgatgtcttt actaacatgg taccacattg cat 353

<210> 30558
 <211> 355
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30558

tattggtggg tttttttcct ttacctgtgt ctcaaagctt ataaaagagc cttctagatc 60
 tgttcttcaa ttgattntgt tcatcaaate ttttttttct tttttttgct ttttatcata 120
 gtccaactcc ttcacaaata tcaccactt tagcaccgac atctcattga attcccactg 180
 ctgacctctc caacctcttg tgtcccttaa tggcatcgt attcactcct ccctcgaagt 240
 ctcttgctct ttctcctttt ccatcattgg ttggatcttt catgaagcta gtgcgatgga 300
 ttgnggaggg gaggagagtg tgatatagat gaccattgtc atatctaatt tggag 355

<210> 30559
 <211> 218
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30559

ttattgatat cttntatagc gagcacggat tcttgatag acattaatgg ggatcaaagt 60
 ttgaatacat ggagaaatat gcattcttgc ggtgaaacta ctccacacct atgtcacaaa 120
 actttatctt atgagatcgc tctcctataa ctacgatgaa ggaatcaatg attattctca 180
 tttgactttt atcgtaaggc ttaactttta tatatata 218

<210> 30560
 <211> 130
 <212> DNA
 <213> Glycine max

<400> 30560

tgatcatcaa accaccttat cccttgaggt tcttcaaat gtttatgtat atagtgtgtg 60

gaaggtcaaa tggaagggttc tccatcaagt gaaacaacac aaacttatgg aggtcagaaa 120
atgttgatgt 130

<210> 30561
<211> 187
<212> DNA
<213> Glycine max

<400> 30561
taccacttgc acggtgctgg aactacttca catggacttg atggggccta tgcaagttga 60
aagcctagga ggaaagacgt atgcctatgt tgttgcgat gaattctcca gatttacctg 120
tgtcaactat atcagagata aatcatactc ctttgaagtt tcctggagct gatctacaac 180
ttcatag 187

<210> 30562
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30562
agctttaact ttatTTTTtca taagcacttt gtgcttgtag attccaccgc aacgcaccct 60
tcttcaaaca ttcggtcata ggacttgcta tagagctaaa attctggata aagcgtcgat 120
aaaatgatgc aagaccaatg aaagatctca cctccgaaac tgttgtagcg ctcggccaaag 180
tcttgatagc atccactttt gcntgatcaa cggatactcc atcttttagac accacatatn 240
caagaaacac caccctttca accaagatat cacacttttc ccttctccca tagagttggt 300
gtgctcttac ggtctcaa atttgtttca aatgagtgaa atgctcctct atagattngc 360
tatacaccac tatgtcgtca aga 383

<210> 30563
<211> 252
<212> DNA
<213> Glycine max

<400> 30563
ttagatatat gtttatgata atacatgttt actctttttt tcttagcata taacgatact 60

caataagtga cgttgaagat gttatagtat agctctgata tgatattgca aattattcga 120
 gtcgatgtat atatatatgg gttgtgtctt gtaaacattg ctatgacatg ataatatgat 180
 atatgacaat cagtgaagta aacagtgata tgtgagctat gaactgtgta gtcacattcc 240
 ttggaaaatc tt 252

<210> 30564
 <211> 208
 <212> DNA
 <213> Glycine max

<400> 30564

ttatcaagta acacaagttg agttttattc acaacattac agtatatctc tcttatctta 60
 ctgagagtga ttctcctata ttcttgagtg attcaagaac accttggtg tatcaaagga 120
 ctttcacaac tctttgtgtg tagtcctcgc tggaaagagt gattcattgc ttcctttcat 180
 catcaccact tgtctttcaa accacaat 208

<210> 30565
 <211> 303
 <212> DNA
 <213> Glycine max

<400> 30565

ttgctagaga cagtgtcaat gctatgtata tggtttcttt ctttgtggca ccaaaggctt 60
 actcatatca gtgagaatgg gctgaattgt ttagccaaga aggatatgct tctacgattg 120
 aagaatgcaa atttagagaa atagtctcat tgcattggtg gtaagaaaac caaagtatcc 180
 ttcaagaaga atcctccctc cagaaaatct gagttgcttg aatcggtgca ttcagatgta 240
 tgtgaccctt tgaaggtgaa atccttttagt ggtgcacttt attcttgtac cttcattgat 300
 gac 303

<210> 30566
 <211> 365
 <212> DNA
 <213> Glycine max

<400> 30566

gaccgggat cttatgtcgc cgcagctggtt acttacattg accctgcacg tgctggaact 60

ctttaatggc ttgatggcct atgcaattga agccttgagg aaagagttgc cttgttgtgc 120
 ggatgattct ccaataacct ggtcaactta tctagaaaat cagacccttg aagtttcagg 180
 actggtctaa acttcaagaa aaaaactgtg tcatcagaga tcatgagtgc catggctaga 240
 gttgaaaaca caattactga atttgccatt gaagcatact catgagttct gcgtatacac 300
 acacaaatgg ctattgaagg aaacagactt gcagagctgt aggcattgctt atgcaagact 360
 tcctt 365

<210> 30567
 <211> 323
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30567

nntgtgtggg agattcagat aatcatctaa agattcatag agtggcttgg aaggagctat 60
 gtaagcccaa gggcaaggga ggcttngct ttaaagaatc cataaatntc aatttagcct 120
 tcttgatgaa gagcgggtgg agcttatgtt cgaatangat gcgttgtggg tcagaataat 180
 cagagaaaag tatcactgtg gagaatcttt gatcccagat attgattgta ataggtttga 240
 gactaatttc tgggtggggcc tttgtaaaac ctggcctgag gtacagaaaa acctttgctg 300
 aaatatttgg gatgggaaca atg 323

<210> 30568
 <211> 303
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30568

gatgctcana aacatcaata tctatcactc catcagtagg tctgcccaga tatttgtaa 60
 tcacagcatg ggagaattta acacactntc ctctgacaaa caccttntga tactcatcac 120
 tttttctgtt agatatgtca gagggaatgt tgacaatgaa ttccttgact aagccttcat 180
 agcaatctcc caacttgctg acagtcttca gcagtccagc agccttgatg aggtccatga 240
 tctccttgca atccaatgca gctctttcca gttctctttc caaggccagt ctctgattgat 300
 aca 303

<210> 30569
 <211> 276
 <212> DNA
 <213> Glycine max

<400> 30569

tgtctcagtg gttatgcgag accgagacca acatgtagc tcttatcagc aagtaccaag 60
 aagaattaaa tctagccacg gcccacgagc acaaagtggc ggacgaatat gcccagagttt 120
 ggttttttag gaaaaacgcc ataactaagc gcaccccaag gcatccttat cgcaccagat 180
 ccaaattctag gacgatgggt gaccaagagg aagtacagga acagatgaaa gccgacatgt 240
 cggcttatat agagcaaagt tcttccatga tggatg 276

<210> 30570
 <211> 309
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30570

aaaaanncga cccggtcctt aagtgacgcg gctgcacttt attctctctg aagactgata 60
 ctgcttttaa gcatatatat actccaatgg tccatgcctt actgctaaat gtcaaaaaga 120
 ggcaaccttt aaattaccca cggggaagat ttggggccaa tagctcctcc tggatgaagaa 180
 atgatcccat gatgaacact tgcaggcttc ttgaaacata gtattgcgcc actccgcctt 240
 gttgcggaaa ataggagggtg aaaagaatca tacttctcca ggcattgcaa atcggcacag 300
 gcaaagaga 309

<210> 30571
 <211> 268
 <212> DNA
 <213> Glycine max

<400> 30571

ttacgtgtat gattctcttc atagagtttg gccctcgaaa ttgttatgag gaggcctgta 60
 tatattaatt gatgcttagt taattaagac tatatcgaat agttgttacc gtgataatat 120
 cattgcataa ttttgatttc tattcttcca atacaaaatg gaaggttgag catggattat 180
 tttaaaacat tgtcaagtag gattgcagca atctcaatta tattattgtc atatgaaaag 240

acattgagac atgggaaaaa ttttatgc

268

<210> 30572
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30572

aacctgtact gtgagagcaa gcgcgagagc tcgacgacaa ccatgactac acggtgtaaa 60
cttgaaggga gggggcagtg ccatgtggcg accggtatgc gaaaagtga gtggaggtct 120
caagatgaaa gaaagtggaa actaagtggg gcggaagtta actanacgca atacaatatt 180
taaactacac atataatagt aactttttta tgaaaaaata ttatntaggg tatttgctta 240
atctacttta tcgtgaaaga aagtatgcaa catattatca aagttttaat ctagtgccgt 300
aaaatactta ggcttttcta aacactataa aaaac 335

<210> 30573
<211> 399
<212> DNA
<213> Glycine max

<400> 30573

atgggtaccc atcagatgtg gtactaggtg gttgtttgtc gatggtgcaa aacaattctc 60
cacatgcaca aatcacgtat aaacccacca tcccctgttg cccacctcaa ctgagctcat 120
gtactccac gtagccctta tcctcgttcc tetcaacgcc ggggtcccat caatcctccc 180
aagcttcac aacatccaag taattcaaca tccaagcctc atgaactaac acagccaaga 240
aaatagggca gaggcagaaa actctgcccc aaacacaaac cgacatcaca gcttttcaca 300
ctcaaatact ccagtaatat tctcttcggtt ccaattcggtt aaccgttgga tcgactctaa 360
aattttactg gaagtctcta gcacataaaa ctacattat 399

<210> 30574
<211> 296
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30574

tggtgcacaa cacgttttcc acatccacaa atcgcgcata aaccaccat ccccttgttc 60
 ccacctgcaa ctgagctcac gtactccac gtagccata tctcgtttc tctcaacacc 120
 gggccccat caatcctccc aatctttccc caacatccaa gtgactcaac attcaaacia 180
 cacataccat cacagccaag aaaacanggg aaaggcagat aattctgccc aaacaccaac 240
 caaaatcaca gcttatctca cttaaaggcc tcagtaacaa nntccttctg ccaatt 296

<210> 30575
 <211> 338
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30575

cctccacggg gtataatgat atcagcgtac ttcttggtgg gaagaataaa gtcatcanaa 60
 gctggcttca caaatntaga atactgcac aagtatatat tataagagac agttactaat 120
 atacgggtaa caaaaaaagg gaattactaa tattggtaga cataaaaaaa acgaaatatg 180
 ttagcttgat ttataatcat aacctcaaaa cagatcattg aagatntaat ntcatanaga 240
 ctgacttctt tacatgtttc taatcaatga aagttcataa cagcctctca gcaacatgaa 300
 ctataacaac atttccaagt cttgaaatgc agtacatg 338

<210> 30576
 <211> 186
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30576

gcattcacc cttccncatt ntcttgacaa accataaata attnttttat agtcgtaacc 60
 ttattgtatt gcaacttaac agcacacaac aatcacttga taaataagtg gcttcagctc 120
 ctatntctta gttttntaa ttaccataac ccacagtga caataatgct aaagcacaat 180
 accata 186

<210> 30577
 <211> 522
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30577

caaaaacggg tggtaagacg tcngcacagc acatcnggga attnagctcg gacccgggat 60
cctctgagtc gacctgcggc atgcaagctt tattgtctga taacgacaaa gagtacactt 120
catcataatt caccatgttt tgtgaggaag cangcattga gcatcaatta acaactcctt 180
acaccctca acaaaatggg gttagtgaag gaaaaaattg aaagataatg gaaatgggtca 240
gatgtatgct tcatgagaaa ggggttaccta acgaatatta ngcagaagct gcgaacactg 300
cagtattcct gctaaatcga cttcccacca aagcagtaaa tatgaagact ccttttaaga 360
cttggatggt ttataaacct tccttgaana atttaaagta tttggatgct tgtgttttac 420
ttatgtgcca cagattaaga gagacaagct agacaagaaa gctgaacatg gtatctttgt 480
gggatatagt tcagtatcta aagcttatag aagtttccaa cn 522

<210> 30578
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30578

ccttcctggc ttgttgtgtt cataatagga ttccttctct tgatgggtgc ttttgtaaag 60
gacatagact ttcagagttt cttgcctaa ggatgtgtga tgcttcacat ttccattgca 120
gtgtggagat tcttctttga gaggaagctt gnggatcttg cacatgaatg gcctangcat 180
gttgttgggg acatagcatt ggcaatatca tgggtctttt ttcttgtgta cacatggaga 240
gagaaatatg attagttaat ntactttata attgtgtaag ttttttgtgc ttgtgggtgt 300
agaaacaaca t 311

<210> 30579
<211> 389
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30579

tagctttgat gtgtgcgtac ccaccatctt ttcatagtag agtatcgata atgtgtctac 60

catcacgatt atcgtctccc ttccatcat tgggggtacc acttgggccg ccagatccgt 120
ccaccttttg ggcgtgttct ttgaaagatc cgtccccctt tgtgcacatg ttctgtagtt 180
gcatcatatc cggaaccata tcacaattgt actgatactg gctaacaaaag gcaaccatta 240
agtccttcca agactggact cggaaggat ccaagttagc gtaccangta acagctaccc 300
cagtaagact ttcttgaag gaatgtatca gacattctc atcttttgcg tattcccga 360
tcttctgaca atacatcttt agatgggtc 389

<210> 30580
<211> 415
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30580

agcacggctg ggnnacatga tatatgtcan gggtttgggt ttattttaaa gggaataaaa 60
nggggaattg tatcatgaga aatgtctgcg gggggtntg aaccatnng cgggcgccga 120
agttgacagc gtgggcattc tccctcctta cnntctttgc accagttgct tccaattctt 180
tttagcattt tggcacttgt ggagggaaaa cgtaatcgaa cttccctctt ttcaaccata 240
cttcaattct ttcctcggcg aatacttgggt ccgcgaagct ggacggcatg taacctacca 300
acttctcata gtaaaacact ggcaagggtgt ctaccatcat cgtgatcatc tccctttcga 360
ccatgggagg ggccacttgt gctaccaggt cactccatcg ctgtgcgtat tcttt 415

<210> 30581
<211> 416
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30581

gatattaaaa attgactata atcagttgag tggcagtaca ncgaaattat acaccaatta 60
ttacaagcca aagtctctga acattttttg ttgcttcttt ttcattcttt catgatattg 120
gatcaaatg gtccccact tcaatatttt caattcgaga ttgactcttc tcatttatag 180
caaaagatcc aacatgacac ttttgctttc acgtacgaaa agcgaaatgc tgatggcctc 240
tatcataaat acacactact ctatacaaac aatgtgtaaa acttcacctt agattttcat 300

gtaactatgc caaaatattg caccgcgcaa aaacttataa cgtttccatt tcataacata 360
 tgtcaggact accgagacca tcatacaact tctattttcc acaaagcaaa tattga 416

<210> 30582
 <211> 332
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30582

ntattcaaga caaagcaatt aaagatattc aagatggatg atctagacaa tctctagagt 60
 cttagaaagg gtatattaaa taggaaggga attccaattg aagtagcaaa tttggcctgc 120
 ataacatatt attatgtaac atttagtgca tgtcaacatt ntcaagtgtt aataacagaa 180
 aattaaatac aactccctgt taagtcgact taccaaaatg tcatcccata ctangtcctt 240
 tagagattct gacacatctt ttcaatttct atgtagcata ggaatctttt ctogaactac 300
 gacccccaaa tagttgtgga acttttattt tt 332

<210> 30583
 <211> 392
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30583

tatcttgtat gtcttggatc ttcttcatca atgaattcct ttgcttcttg aggtttgatt 60
 gcagcgaagt ggagaaggag aaagatgaat ggagatgccca cttcaagtag aagatgagtc 120
 tagaagaagt tcaccaccat aggaagccat ggataagagc ttgaaggtag agaagatga 180
 atgaaggag aggaagagaa gagcatgaaa tttagtgcct cttagaaga ctgaactttg 240
 aagtttaatt ctcatatgat caaagttgaa aaaatgcaca cacaagacct ctatttatag 300
 cctaagtgtc acacaaaatt ggaggggaaat ttgaatttct attcanattt cactcgaatn 360
 tgtggagcca anatatcact aattatgatt ag 392

<210> 30584
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30584

tcatgatgaa tcaagagtga ttcanagatg ttttgatgat atctaagatg ataacaaaag 60
atgatgacaa aggtgatgac aaaaagctca aaggtcaatt aaagaatgag ttcaagatat 120
tcaagataga atcaagaaca cttcaagatt caagaggaaa gttgatttca agaatacaaga 180
gatcaagatt tcaagaatca agattttaagt gatcaagatt caagactcaa gattcaagaa 240
tcaagagaag acttactcaa gataagtatg aaaagggttt tctcaaaaat tgagtagcac 300
atgcattttt ctcaaaacat gtttaccaaa gagttttact ctctggtaat cgattactag 360
attgttgtat ccgataccag tagcaaaatg gttttgaaaa aaaaatcaaa tgaatta 417

<210> 30585
<211> 418
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30585

agcttgtcat attatctctt caccaacctt tggtcccacg atggattcat atcatcttca 60
tatgactgga tgaatcttgt aatatttgga ttaacaacat taaaattctg acacagaaat 120
acacaaattg agaaatgaga tgattatgca ggaaggcaaa acttttttcc caataataat 180
aaagaaattc atattcacia tagaagctta aaaaatatct taagtagggg aaaaaatnta 240
ctcatgttca caatactttc aatagagtta aacatctagc atatgataga agctcatttg 300
ctattgctag gaaaaggctt atctcattat ggcgaggcaa caagctccag gaatataaga 360
aaaggaaata aactctcacc agtttttagct tacatanngg gaacataaaa gcttacat 418

<210> 30586
<211> 307
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30586

gcttgtacaa gcacataaag ganaagggaa actgatgaat gtgttttacac atcctttgca 60
caaaagatta ataggcctaa ctatctaaaa acagtcccca gtggagttgc caattgtcac 120
aacctaccct ttggcgggtg atgtaagctc cattggagct tgtaggccta ggatcttctt 180

cattaatgga ttccttcgct tcttgaaga tgaatggcag cgaaatggag aaggaataga 240
gagaggagat gccacttcac tgaaaagatg agtctacaag aacctcacca ccatangagg 300
caatgga 307

<210> 30587
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30587

tttagatcag aacttctaaa tttaggggtct ttggaatccg gacgcgaaca caaatatatg 60
caagtaacca tagatattta agaataatat ttgttaaaat acattntctt aatcgataat 120
aaataaataa ataatatattt taacatatat ttatatataa aatgaataat ttgaaaatat 180
atatattcat actaaaaatt aattaaacca acttatttat atataattag agacatttgt 240
ttatgcaggat atccgttaaa ctgttcaaat ccaatataaa caggatntat ccattntact 300
caaccaacat gcacccatgt tctggatcat gacgggtttg cctgagtgca gtctaaaaca 360
aaagtcattgt gattnttcat ttat 384

<210> 30588
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30588

ttgcacatta nactccacat ttatctctat agaaaaaat atattaatac ggaaccctag 60
ctacttttaa aaagacgcaa caagggaatg tttntctctt ctgttcttta caaatagcta 120
ggataatgat ttttgtacaa ttattctgtg tataacagtt cctttgattt ttgtgaaata 180
tttttttgat atacatgttt atgaaaaaat aaaatttaag tgagaaataa aaaaaaatc 240
atatatgata atgagatagg atcgaagtnt anaatttgat gaaaccttaa ccgcagtgtt 300
agtagtaaaa agaattctta cacatgcaac aanagtcgc taactgctaa taaatatatc 360
gatggtgaca agacatagag tacangcttt ggggttggtta gttaaaacat gttcacatg 420
acgtagtaat tacaaaaa 438

<210> 30589
 <211> 412
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30589

caagctgnngg tctanntcta aggattgagc atggcatagg aagggttag tatgttgaat 60
 ggtangaaaa tttgataatg atcacctcca cagactctnt gacgctcaac tttggaataa 120
 gagaaataaa ataaaaagtg aagattaaga agttcatata taaagggtaa tacatctcta 180
 tatagtgatg attttggcgg aaaaaatatt atgtactctg agagcatgtg acctacgaag 240
 cttattaata aggaggaaat ccatgcaatc tttgtgatat agagtagaaa gtacatttaa 300
 gaatgtgttg ctgaacttgc tcataattga atgtagagtt aacaattcag gtgacaagta 360
 taccaagtaa tgacattatc tatctggatt ctggagaatg aatatgagca tg 412

<210> 30590
 <211> 279
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30590

tagganaatg atcttagagg atttatatat gagcaacttg tgtactccac cctgggctgc 60
 caagatgtat caggatctca cgacaatggt ttggtgacca aacatgaaga gagaggctag 120
 tgagtttgtg tatgtgtgta tagtatgtca gatcgctaac atagaacatc tgagaccctc 180
 atgtaagttg caacactttg agatacccag aggaagtgga atatttttca ttgatttcat 240
 tgttgactac ctaggacccc caaggtttcg atctatcta 279

<210> 30591
 <211> 270
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30591

acttcctca cgtactgtct cgtgggggggt atgaactgct tgacaagaaa cttatggagg 60

agaagagcaa gcgtggacat gaggaacatt cgtgtactga aagcccaaca ctcaacgtcg 120
 acccaccatc cccagttgca agacacttga agtgggaagat cgcccgact aagcggcatg 180
 gccaaatgac gtctgaagtgc gcacaagaaa ttgtagacan aatgggtcagg tcatatattt 240
 ttttggttac tgtcattggc anataatggt 270

<210> 30592
 <211> 401
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30592

gcttgtgatt atcatgtaga ggattcatca naaggagttg accacgaaca aactataag 60
 attcattcaa ccccatgaga aattgcatca acttgctct tttgttgttt gcttcacatg 120
 tgcaaatagc atcattatag gatcctaact catccacaa tttttttta actttgtata 180
 gtatgcagca actatcattt gatcttaagt aaggcaagca atctctctct cgatctggaa 240
 aatgagtggg gcgttgcttt gaaagaaaca attttgaaga tcttcccaa cttcatgagc 300
 agtgtaaana aaaataacac tatctgtaat atcaggggtc attgaattaa gaatcccatg 360
 acaaaccata tcattgcatt nttcatgcta catagtcttc t 401

<210> 30593
 <211> 331
 <212> DNA
 <213> Glycine max

<400> 30593

tgatgggtac catgaggtgt ttttgtgttt gaccacgcg ggtgttgaag agacggcatg 60
 ggcattcctt tcttctcttt ttgcccctgt tgcccgcatt cttttggcgt tcacgtttgt 120
 ggaggaaacg taatcaaact ttctctcttt caatccaacc tcgattcttt ccccgcaaa 180
 caccagatcc gcgaaactgg acggcatgta acccagtatc ttctcatagt aaaacactgg 240
 cagagtgtct accatcatgg tgaacatctc tctctcaacc atggaggagc tacctgtgcc 300
 gccaatccct ccatcgctgc gcatattctt t 331

<210> 30594
 <211> 426

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30594

 agcttgtcac ttgtcagccg aatggagact tctgactaac gatccttaga tgacaaggta 60
 ctaccacatg gctacttggc ttanagaatt ctctcgccaat tttgagctca gacacattat 120
 aagagaaaac aatgagagcc aacttggtgt taaaactcgc cagtacaaag aagaaagggt 180
 aatacataat agttatcaag gaaacaatct tggaactagg cttggacaag gtggtagcga 240
 gtgtaactct catttattga gcggtggataa ttgagattta cgacttcctg gaaataactc 300
 actctcgaat aatctagtag cgacaagaaa gattaagaga aatgccagtt attatgtgat 360
 agcggggagga tacctataca aaagacgctn tacaacctct ctgttgaaat gctaagtcgg 420
 gatcat 426

<210> 30595
 <211> 213
 <212> DNA
 <213> Glycine max

 <400> 30595

 ttaccatcta tcccagccct ctccaagaat aacttgtcat gtaaagaaag cggtcctata 60
 caccaaagt ttgttaccac tctctcagac cctattacat acatcttata cgactgctga 120
 aaagtgatca ccaatatccc tgcattgcaaa tttccccagc tttgaattgt tcaacggagt 180
 gcatccaaac ctgtcagtcg tgactcattg ccc 213

<210> 30596
 <211> 503
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30596

 agaacgaagg ggggtggaac gtacgcagca tgcgnncata nagnncgncc ngggatnntt 60
 nagtcacctg cggtcatgcaa gctttcttta ttgcgtcacg tntannaccg agctcgatgg 120
 tgggtgcgagc ctttgatggt actcngcggg aagtgatggg ggaaatcgat attcccatc 180
 agatacgccc ccacacttgc catgtggtgt ttcaagtaat ggatataaat tccgcctata 240

agctgctctt gagaagacct tggattcatg cnctgngagt ggtcccttca acgctttacc 300
 agaaattaaa gttcgcagtg ggtggacttt tagtgatagt gtcnggtgaa gacgatatgt 360
 tagtgagctt ccactcctcc tcaccgtaca tagacgtggc ggagaaatca ttgaaccggc 420
 ttcttatect ttgnggggtg agctgtgcct cngtgggaacc agtccgctct acctttctct 480
 ccacgccgca taatggtgca ccg 503

<210> 30597
 <211> 136
 <212> DNA
 <213> Glycine max

<400> 30597

aaaagctgca tctacagcgt agaagtcacc tgcagttatg ttagctagca catccatttc 60
 tagtctgtat cttgttggtc atcaactagc atgcagtcct gaaatattat agaaatgaca 120
 tatcttatgg aagaaa 136

<210> 30598
 <211> 316
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30598

gcttccttgg gaattaanat gattaaagct tccttaagaa gctagagggg gactactcat 60
 atccctccaa tagctaagtt cacacctatg ccaaaataca tgaaaataca atgggaagca 120
 aggaaggtag cttccttggg aagcaaggaa gaaagcttcc ttgagaagct agaggggggg 180
 cgggtggctac tcacaccgc tcaatagtta tgctcacc cc catgccaaaa tacatgataa 240
 tacaanaaaa taaaagtccc tactacaaag actactcaa atgccctaaa atataaggct 300
 aaaaccctat actact 316

<210> 30599
 <211> 503
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30599

aagcgtttca taccanngct tgcacnnncn nnnannaaga ancnngggatc cttanagtcg 60
 agctgatagc tgcagcttgt atttgttttc ttttaatgtg actatgaggc aggcgcgtct 120
 ttgagaacca atccatcgac ctctccatct caaaactcaa taagatataa gctccacaca 180
 tctcagctca aaccaataa atggagctag tatgatagat agaacggctt gctataaact 240
 gcctgccaca caagtaatcg acattataga tgggaaatct gtgacaactt ctttctgtca 300
 gagcaattgg agaccactta attgaacca ctgtatgcta tcgccactac attaaactttt 360
 ttctaaggat caagtatagg tataaatata tatgtattaa ctttatgata actaagtttc 420
 atccacacaa ccagcttggt caacctttag tgaaaatata cttctcaact atgtatttta 480
 tatctatata accaaaccac aag 503

<210> 30600
 <211> 379
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30600

cactatanaa atactcccaa gcttgtaggg attanntggg gtacccatca caatgtggta 60
 ctagtgtggt ctgtcgggcg aatgggtgcaa aacgattctc cacatccata aatcacgtac 120
 aaaccaccca tcccctgttg cccacctcca attgagctca cgtactccca cgtagccctt 180
 atcctcgttc atctcaacgc cgagtcccca tcaatcctcc caagctccca caacatccaa 240
 ttaattccac atccaatcat catggactaa caaaaccaag caaaacaagg caaaggcaga 300
 aaactctgcc caaaacacaa ccanaatca cagcttttca catacaaata cccagtaac 360
 atttccttcg ttcgaattc 379

<210> 30601
 <211> 409
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30601

catgtttgct ttgcttctgt gatgatgtgg caggagcaat ttctgatggt tttaaaacaa 60
 cagcattacc agctgcaata gctccaacga ctggatcaag tgacaacact ggaaagatag 120

acagagaatg taatatagat nttgcagaga ttggaggaaa gaaaaagggtg agatgtcagc 180
tcaaaagtat gtgacacaat accaactaca tattaacata cagaatgggt agttccatgc 240
agagatgact aacacaaccc ccagtgggtc agatactatt tcagctgaag atggaaatag 300
tgcgattgaa gtcttgacct gcaataaaag ggaaattcat gtgataaatc ataagtacca 360
aacaatgtca gcaatgaaat tcacataatg tatanccttt caggagtca 409

<210> 30602
<211> 318
<212> DNA
<213> Glycine max

<400> 30602

caacatatat atgttcatcc attctaagct atcctttttt ttttacatat gttgtaccag 60
gcctcccatc cctctaaaag tgaccctaatt tctcatttta atactaatgt tgccccataa 120
ggaaatgcc aatatgtttc ccagagataa aagaaacatg tctctcaaag ttctttgtat 180
ttctatatat ggccatattc tctaaatata ttaatagatt atttactgag agaactcaag 240
attcctagtt taaaaaagaa gtcacaccat tagccctctg ttttattctt ctctatgatg 300
tcgtttgat tcttctct 318

<210> 30603
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30603

agctntgagc ttattcttac gacaataact nttgattcgg atgtccgatt gtgtcccgt 60
ttatatcgag atgctcgtaa ttgaaaatag aagctctgag caaattcaaa cgacaataac 120
ttttgactcg ggtgtccgat tgtgtcccgat agtatatcga gacgctcgaa attgaaaaca 180
gaagcactga gcaaattcaa acgacaataa ctttttactc ggatgaccga ttgagtcccg 240
taatatatcg agacgctcgt aattgaaaac agaagctctg agcaaattca aacgacaata 300
acttttgact cggatgtccg attgagtccc gtaatatatc gagacgctcg caattgtaaa 360
cagaagctct gagcaaattc acacgacaat tactttctac tcggatgtcc gattgagtc 419

<210> 30604
 <211> 419
 <212> DNA
 <213> Glycine max

<400> 30604

gacacataga aactcaagct gttcttgatt tttctaagtt ctttaacaag cttagatcaa 60
 tataacttgct cttcatttaa ttgtctttgg gcttggcggc ctgatcaac aaagtacttt 120
 cggcacctac tatatgttga cttgaccaac gctcttatcg gtatgctgcg acaatccttc 180
 aacaccttat tcacacattc tgagagggtt gttgtcatgt gaccatatct tcgtccagat 240
 gtatcataag ccatggctcc attttccttt gaaatgcgat caatccatgt cgctatggct 300
 ggactcaatt gacaaaattt ttctaagttt tgatcaaaca catgcttgca aagagtgtac 360
 gctacatcac aattgttacc atcaaaagtt gaggtagata tgaaactcaa ataacttca 419

<210> 30605
 <211> 423
 <212> DNA
 <213> Glycine max

<400> 30605

ttcttgctca tttgtttaat aaacttaaga tcaatagtta ttataaaaaa ttctatcaat 60
 aaaatttctca catcacacta aaaatcaaat ttgcatcctt ataaaatatg atttctatcc 120
 ttataagtgg actcacacct ttgttggtta tctcactc taaatatgta tatgtattgg 180
 ttctgtcatc aacaataacc ctttgcaatg tcaagtaaga ttgtttctac taaaaaata 240
 tgtgatatgt ggcatagtaa taaaacatat tgattaatga agatcattat aaatttagta 300
 taaactgata ctggaaatga ttacaagcgt gggaagagaa aaagagtgat gaggtagtgc 360
 ttctgcaata atttaccac aggagcagtt ggcagtagtt agtgcgtctt ttaaatgata 420
 cat 423

<210> 30606
 <211> 339
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30606

tatcctctac gacaatcaac tcggcggcaa aattcccggc tctatcggga accttaagag 60
 ccttcaagtg ataagagcag gtggaaacaa gaacctggaa ggccttttac cacaagaaat 120
 tggcaattgt tccagtttgg tcatgttggg tcttgctgaa actagccttt caggttctct 180
 acctccaact cttggcctct tgaaaaacct tgaaaccatt gccatttaca ctccctact 240
 ctcaggtgaa ataccacctg aacttgngta ctgcacaagg cttccaaaca tatatcttta 300
 cgagaactcc ctcaactggat ccataccaag caagttggg 339

<210> 30607
 <211> 424
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30607

agcttgaact tgctgtatct tctacttttc tagtgatttc atatgagaca ttgtgttcat 60
 tctgttttat ctgaatataa tttagtatat acacaaatat tgattttgat atgctaataga 120
 gatatgctat ttgcatatct agaatccata tggcagtttc cttgacgagg actataactg 180
 aaagccaaaa cgtgtttccc atgtctatct gtcttgcaag gcggatttct gatcatggag 240
 cttcaagggg agggatactg cttcttaaac gttgattcaa ttattattcc ttactccttg 300
 gtgtaaaata ccttatatcn taataaattt tatgctgggt tatactttat gtnttaactt 360
 gtcaaaacaa aagcaaaccg ccattgatcg aattggcggg gatttcatct tccgaaactc 420
 ccct 424

<210> 30608
 <211> 502
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30608

ggggnnnnnn nnnnttggga cagtcagact cngcaatcac tcgaccgggg tctctgagtc 60
 acctgaggca tgcaagctct ntgtataatc tttcttggaa agctagagct agctacacac 120
 accctctaa taactaagct cacctncttg agaagcntnc ttgagaagat tccatagagaa 180
 gctagagctt aactacacac acctctctaa tagctaagct cacctncttg acatgagaag 240

ctagagctta gctacatacc cncataata actaagtnta accccatgcc aaaatacatg 300
 anaatataan aaaagtcctt nactacaaga ctactcanaa tgccttgaaa tacaagacta 360
 anaccctata ctactagaan tggcaaaaata cannggccan aaaanggana acctattcta 420
 tatttataaa gngagtgacc caaccttgct catgggctag aatctacctg tgtcatgaga 480
 cccagggcct cttagcactc tn 502

<210> 30609
 <211> 387
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30609

aatactagct taggttttcc tgctttttta ttctnctacc tcttttcttt cttcccttag 60
 aaacctttct taagtttaaa atcctgcgac tacgcgctaa gcgcgtaagc tagctaagcg 120
 acccatgtgc gctaagcacc ttttcacttg actgcaggtg ctctatgctg cttcttgcgc 180
 ctgagcggac accctccac taagcaacaa tagctcgcta agcaagtccg acgcactaaa 240
 caciaaacat catgcttcaa cttctctctt tatcccttgc ttggatatct acaaaataaa 300
 atcatcaaac agtttgaatt aacgatttaa ggtacctact gcgcaaatac ttcgaggata 360
 ttaaaattat aatgattcac acaaaaa 387

<210> 30610
 <211> 154
 <212> DNA
 <213> Glycine max

<400> 30610

tctgggtggga catcttgact tgctttctaa tctgacattc atttcagatt ctgccttctt 60
 ctatgttcag attgggaatg cctctagcag cacctttgtc gatgattttc ttcattgcctc 120
 ttaagtgcag atgtgcaaat ctttgatgcc atat 154

<210> 30611
 <211> 398
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30611

catgtntgtc atcatcaaaa atgCGGagaa tgtgaatgta ttttgatgat gtcaaaagaa 60
gaatcaaacc aggctcattn tgcttcaa atacaaga ttgttcaaca aacanagcct 120
tgattcaaga tttcttcaag atcaagtcgc gcctcacaat gaaagggttc aagtcattca 180
aggcacatgt aatcgattac caatggtttg aaagtgtgtg atcnattaca catcatantg 240
tatcgactac tacagactct gaatgtggga attcannatt taatgaaggg cataactgtt 300
cangaaaata actgtcgtat tattacacta anntctgtat cgatttcaga gaggatntca 360
cggatatcgc cacagcacat ctatcattcg attttgag 398

<210> 30612
<211> 242
<212> DNA
<213> Glycine max

<400> 30612

atgacttgat attgactttt cgggaatgca cagacatctc acattcatca aactgggtcca 60
gtgaaggatg tttgggaaca gttgccggtc tctcataatg agccgatgat gttcatcatc 120
ttggctgacc ggttgcgcac tgctggggat aatgtgaaaa ttaccacccc ttgtctaagt 180
ccactctttg tcaccggcta atgggttatac atggacattt tcttgcccac ttgacagtc 240
tc 242

<210> 30613
<211> 241
<212> DNA
<213> Glycine max

<400> 30613

tatgcagtta ttccccccaa ccacaagaaa gcaaaaccct taaattccat acgaatcaaa 60
atcctcaaca gagttttacaa atccgaaacg aatagaagaa attggaatta aaagaaaaaa 120
acaaattata aaaagaagaa caaactaact aattggatcg tgggtggaac ggtgtgtgat 180
gCGGctgatt attttcgatc ctctgttccg tgcttcagag agaacacaga agaaatgatt 240
t 241

<210> 30614
 <211> 339
 <212> DNA
 <213> Glycine max

<400> 30614

ggattatata tcagagtgtt atgatccagc cttaaccata atcatgacta acataagcta 60
 tgtttaataa gacattctag ttaatttctt atttttatta ggtgaaaaac tcaattgtgc 120
 aataaacaag ttcttataat acctactaat gcttgatatt ctttttaatg atacattttt 180
 taagtacttt aacatctaata tttacttaac aattctaata taactaatta gtctttaact 240
 gcttactatc aattagttgt tttaacttct acctaacctt gtaacttcta gctaattgtg 300
 caaacgtaat gataatcacc aatttggtat tctcttact 339

<210> 30615
 <211> 488
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30615

nggctgtaag accatggcna gcacatcnng caattcagct cgacccggga nctctgagtc 60
 acctgcggcn tgcagctttc ttgtttgtcc atcnngngaca attaactgta aagtgggccc 120
 aattggattc taatttcaac ttacctatnt ggaagtgaca tcatggcagt taggtcccag 180
 ctntccattg tggattcagt cacanaacca acttcaatat gtnggactat ctaacacggn 240
 gatnttcgat tctattccca cacacgatgt gggaagcacc ttctcanggt ttgtatntaa 300
 acctctctcn taatcatatc catgggtgaga tngnactaca ttaangaatc aatatctatc 360
 caaatattga tctaagctag atcactcgtg tggttaattac cctatctttc acgtgatgtg 420
 ctcnngtaga tctntcangc attcattctc tgatccatga atgacttnta tgtacnatca 480
 ngacanca 488

<210> 30616
 <211> 487
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30616

tgaccggcga anagttgaga ccctggtaga atccgatacc nctgtaanaa cacngcnng 60
 gggatctact acgctttag aactatggat gctgttcaag caaattacac tatcttagag 120
 aaggagctat tatcgatagc ttttgctctt gagaaattac gttcatatct gcttggtact 180
 cgtgttattg tttatactga ccattgcact ctgaagtacc tgttgaagaa cgctgaatca 240
 aagcctaaat tgatcaggtg gatgctttgg atccaagagt ttgatttga gatccgtgat 300
 cagatgggta ccacaaactc tttgggtgac cacctgagta tgattgagcg tgcgcctgat 360
 gactcaccca ttcgggatga attttcacat gaccatttgt acattttgta taagatctct 420
 gattccgtcc ccactccatg gtttgcttat attgcaatta tatggctgct catgttttcc 480
 tccctcn 487

<210> 30617
 <211> 312
 <212> DNA
 <213> Glycine max

<400> 30617
 ccggaatcca ttttcgggga aaaatgtata gactgagaaa agaaaaatca atacggcgaa 60
 tgttttgtgc tagcaaaact atgggacctg ctttgtggta ctactctggt gtgaaatagc 120
 gatttcacat tactgatatt gaatttggct catTTTTTat agacgatcgt cagaactctc 180
 atccttggtc tcctctatTT cctcgaagt atgactctaa tcttgagtct tttcttttgg 240
 tataaactaa tcttgagtct gaatgggtgg ttaagtaaT ttctaattga aatgatactc 300
 taatctaaaa tt 312

<210> 30618
 <211> 356
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30618

aatcacatgt ntgtcatcat caaaaatggg gagaatgtga atgtattttg atgatgtcaa 60
 aagaagaatc aaacaaggct cattntgctt caagattaat acaagattgt ttcaacaaac 120
 aaagccttga ttcaagattt cttcaagatc aagtcttgcc tcacaatgaa aggtttcaag 180

tcattcaagg cacatgtaat cgattaccaa tggtttgaaa gtgtgtaatc gactacacat 240
 catatgtaat cgactactac agactctgaa tgttggaat tcaaaattta aatgaagggt 300
 cataactgtt caagacaaat aactgtgtaa tgcattacac taattctgta atcgat 356

<210> 30619
 <211> 372
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30619

tgagcgaatc caaacgacaa taactgtgta ctcgatgtc ttattgagtc ccgtaataata 60
 tcgacatgct cgaaattgaa tgttgaagct ctgagcacat tcanacgaca ataacttttt 120
 actcgatgt ctgattgagt cccgtaacat atcgagacgc tcgaaattga atgttgaagc 180
 tctcagccaa ttcatacgac aataactttt ttctcgatg tctgattgag tcccgtcata 240
 tatcgagacg ctcgaaattg aatggtaaag ctctgagcca actcatacga caataacggt 300
 ttactcgat gtctgattga gtcccggtac ttatcgagac gctccgaatt gaatgttgaa 360
 gctctcaacc aa 372

<210> 30620
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30620

tgctttgaat attggtcttt gccagtgaat ggatcgatgt gggatatgaa aaaggcaaat 60
 ttagtcatcc tgcttgagc aatgagaaaa ctggggcaaa tgaagagggt gagaaagagg 120
 gagaaacca tgctgtgact gccattccta tacggccaag tttcccacca aaccaacaa 180
 tgtcattact caatcaataa caaacctcct ccttaccac caccagttta tccacaaagg 240
 ccatccctaa atcaaccaca aagcctgtct accgcacttc caatgacgaa gaccaccttt 300
 agcacaacc aaataaaaca ccaaccaaga aatgaattnt gcagcgaana gcctgtatga 360
 ttcaccccaa attccggtgt catatgctaa ctttgctcca tatcta 406

<210> 30621

<211> 422
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30621

 agcttgcttc tacaatctcc ccctttntga tgatgacaac cctgaaatca agaaacacac 60
 acacacacac acttttttct agtcgatcac tcaacttaatt ctccatattc tccccctttg 120
 tttttgagtt tatgcttcac ttgaaattaa gttaattact tatgtgagtt cttgatttaa 180
 tccttatttc tctccccctt tggcatcaac aaaaagccaa agtgcgtaac aaatataaat 240
 catacatata ttactaatca ttcacaagac attcattgaa aaatctaaac caatcatgaa 300
 gcaagaaaca tgaatagatc anatatataa aatccacata gtcataatac acaattcata 360
 attgttcaat catactatgc aaataanaga aaatactaaa tgggtcanatg tcataataat 420
 at 422

<210> 30622
 <211> 390
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30622

 totcaaacat taatntaaac aataaatgca caacttagca tgttttagtgg tgccaatagc 60
 agttaaaatc actaaacaaa ttcacatgaa aggaattgat ttgaactaat gaacaatagc 120
 aaacaaaagg aattaattga tgcacaagca ccaagagcaa tacaaattga tgctaataat 180
 aatgacgcac cacatagaaa ttagaagcaa aaattaggct caaattggaa tatgatgcaa 240
 tcacaaagag aaaggcactg aatgaattca anaaaaaacc gacgcatggg attaaaatgc 300
 tataacaaat taacacaata tgccacaagt aggtgacaca catccaattg atatttgatt 360
 gatgcacttc atacaaccaa tgctaattc 390

<210> 30623
 <211> 361
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30623

agctcgaccc gggctcctcta atcacctgca gctgcagctt caatttcaag cnaacagcta 60
aagaagaaaa tgtgggatcc tgtgnaacca atggtgcttg gaaagaagag ttaaaaactg 120
ccgttaataa ttatgcttct gtcattacag aacttgatgt tgcaaagaaa gaactgagta 180
aaattcgta ggggtatgat ttatcctcgg aagcaagagt ttctgctctc aagcgagcat 240
cagaagctga agatgcaatg aatgcataca ccataagagc atgtgagcta tctaaagaaa 300
ttttggctgt acaggaatca tttgagaaaa cgaatgctga atntgtccaa gcacatcaac 360
t 361

<210> 30624
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30624

agcttgaatt ttatgttgat ttgacagaga gaaatacttg tagcttggtg aagttactag 60
aatttggtgg tttgctataa gaacttgaca ttgtcttggt ggttgagatg aaccaacata 120
aatntgatgt gtcttattct tttttatttc tcttttgcta tttgatctgt tagggtttga 180
atttgatctt tattatttaa aaactttggt tgttttacia agatttgaaa ctatcatctt 240
atttgttntg caaaagtctg atatctgttt tgttaagtct tacttcacaa gacaataact 300
ntattanttt acgaaaaaat tattttttta tgaaaattac aattcaatct tatttcttgt 360
aatatttatt tttgcaatat tattatattg tat 393

<210> 30625
<211> 431
<212> DNA
<213> Glycine max

<400> 30625

tcctcgggtgc cattcactgc gattgctaac atttggaag ctagtttacc aagaaatgct 60
actcttaaaa caaatatggc atacaacctc ctccaataaa cacaacatc aatgtaaatt 120
tagagcaaac tcatgcacat acttccttat gaacattcac tcgcacaaaa tattcttcta 180
cctaaaaaaa atgcacccat gcgcaatcaa agcacctttg ttacctagat atatttatgt 240

gtacttccaa ggtgtattta ctacttacat cacatgcatt tccttggcta aatgtacata 300
catgcatact caaagcatct tggctaccaa aaattgcaca cgtgcacatt ctggcatttc 360
tagtacctat gcatatacaa actatgtgat gaatctatgc tatctacaca ataaggtgct 420
acatttcatg c 431

<210> 30626
<211> 381
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30626

ntagcttcca ctcgtaagac catatgccac tagaccagtc tattctagaa tatagaaaat 60
aaacttcatg gtctcgaata catgtagcgt gcctaaactn tgaggtgtaa catcccaaaa 120
ataatctaata aattatctag ataaagatat ttaaataatat cttttattac acgacaagat 180
agattaaata atttaaataat atcatataat gatttttatt tcgaacagag gttactataa 240
taatataata gatttggttaa cgataaatta acgataatag aaagtataga taaagaagga 300
tcatttaaca gattgcaaata tacacacaat tcttaataata tgggtgacag tctttctctt 360
agaacacaca cataaaggat c 381

<210> 30627
<211> 403
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30627

tcacttgcca ctggttataa actacaacct ttaaacaaca acttgcaagc aattaaaatt 60
cgtgactttc taagagacaa ttgtatcttg actccacgat ctagctcttg aatcatctta 120
cacacccttt gtaacctaaa gtacaagaat tccctacctt cccctcatt gcattattac 180
atcgtcaagc acattctata taagttatga ccctatgtat catctaaacc ctgtaggaga 240
tgtgagaaga aatcatata taaacatacg ttttaataaac ctatataaca ctgatngaga 300
tattcttacg gatgcttatc tggatatctt agaattacat tgcttcggtg cttgaaacgc 360
caaccaata tccacattca tatctctcac tgaaatagga cgc 403

<210> 30628
 <211> 370
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30628

accaagtgtg tctaactttt cctaagtcta tgtttttgcg ttgttgcgct aaacgcacct 60
 tgcgcactaa nggagtactg ttatttttat aaggcacgct aagcgagcca gtctcgctaa 120
 gcgcccattc tatttttttag ttttattttt ctgctttcag ttaaaataaa agcatgtcta 180
 atatgattat tgtgcttatt ttttatgcag atgacctcca ngaagaggaa agccatagcc 240
 tcccgatccc gggaaccata taacaccacc cattntgttt ntgaggtcgc ttangagcga 300
 tattctcaaa acattcacac caggaacatc cttccagaga ggaatgttaa tctttttgtg 360
 atagagtatg 370

<210> 30629
 <211> 323
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30629

gcagctttac attaatcca gctctcgatt gtacgggctg atcagacatc gagtaaaagt 60
 attgtcgttt aatttgetca gtgcataaca ttcaatttca gcatctcgat acgtgatggg 120
 actgaatcag acatttgagt aaaagttatt gcgtttgaat tgctcagtga tgaacattca 180
 ttcagcgttc gattataccg actcatanac atcgagtana agatttgctg tgatnactta 240
 agctcacatt cattcacacc taattgtacg gactgatcga catcgagtaa agtattgcgt 300
 tgattgetca actcacattc att 323

<210> 30630
 <211> 250
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30630

tctttgagct tcattgttgt cctttaatgg cgattntcca ccatggagat gcatcggaag 60

acaaaggaga ataggtaaga ggcgacgcca tccactatgg aataagccat gaatgagctt 120
caccacgaag atgagcaaag agagtgttgg atcgagtggc ctcanaatca ttaagaaggc 180
gggggggggtg aattaattat tcctaaacct ttactaatta aaaaattact cttctaagtc 240
ttttacttat 250

<210> 30631
<211> 366
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30631

cccacaacaa caacaacaat tcatcaaccc acgatggaga agaagaagcg gtgaaacccg 60
accagctgtt caaggaaacc gcagagtaca tcgtgttgct gcggacgcgc gtcgtggttc 120
tccagaaact cattgagtat tatggaaaca acaacgacac cacccaagat gagaatgaac 180
atgaagatgg tgtcttgttt acatagctnt ttcactctct tcttcttctt catcttatta 240
ttattattat tcttttttgt tttcttctta catggtttgt tttgtgactt ttgtcctttc 300
attaatgaag aaaaaaacaa aagacaagat ctttgggtcta gtgttttttt tcttggaggg 360
ggggggg 366

<210> 30632
<211> 369
<212> DNA
<213> Glycine max

<400> 30632

ccaaaatcat gagcacgaga agagtctgtt ttcttattaa tacaattcat tgtatatgcc 60
gtgatcactc aacattccag tcttatgtaa gaacatatta tggattaacg gacaataata 120
cctctatgtc tgagatcgct ctacacaaat aggctcagca tgtgggacgca tagaaagtgg 180
atagtggcaa aactgaatac atgctgctat atacaattaa tagactccca attgtctatt 240
atttgaataa gacctttgat ctttttctaa aataagcgga gctaaaagat tatctgtttt 300
tgatgtcaat attgaaaata gtactcttgg ctgataaaag tgaaaaatgc ataatgccgg 360
ataaaatat 369

<210> 30633
 <211> 346
 <212> DNA
 <213> Glycine max

<400> 30633

ttcgaatggt cttatacttt tcaactcggat gtccgattcg cgggcataac tcatctagat 60
 gctcgaaatt gaacatcgga agctctcgag aaattcaaatt ggtcataact tttcacacgg 120
 atgtccaaat ttaggacata atatatcgag aactcgaata ttccacaacg gatgtactcg 180
 agaaatttga atggtcataa cttttcacac ggatgtccga atgtgggaca taatatatcg 240
 agacgctcga aattgcgcta cggaagcact cgagatttcg aatggtcata acttttcaca 300
 cgaatgtctg attcgcggac ataactcatc tagacgctcg aaattg 346

<210> 30634
 <211> 380
 <212> DNA
 <213> Glycine max

<400> 30634

agtttcaaca taataaagac tttctcaagt caagaactat gatcttcttg aggaactcta 60
 ggtttcaaatt cgtgccatag taagcatggg atatattgaa taacaattac agggagaacc 120
 catgcattga tcttaatgat gtgtgtaatg agacaatgaa tgatgcacaa ttcgaaagct 180
 ttgttgatat tttgtcacc cttggaaatt gttgctcttt tagtctactg gcagttgaat 240
 cttttccctc tgcttcaatg ccctctcagt caatggcatt ggcttcaact aggacttcta 300
 caccatgtgg gacaaagtgt aaagttgtta tgggtgatgt cactgatata caatttgatg 360
 agttcaacac aggtcttgat 380

<210> 30635
 <211> 316
 <212> DNA
 <213> Glycine max

<400> 30635

tatgaagaga gaacaattta gagagtgatc gaagactttc aaatggattt ccaactgaatt 60
 tattcataga gagatcgaga tatcttaatg atgaaagttt tccaaatgat ctaggaagag 120

caccaccaat tgagttgttg gaaaaaagta acgtgtcaat atttttaaat gcccgaatat 180
gatctgtcag attgcctgaa agtcgtgaac tctgaactgc aagtcttgtg agtccatggg 240
aaatacaagg agcaagaatt tctaaaagtt cattaacctg ttggttgagt ttgagatatg 300
ataaatctat caccct 316

<210> 30636
<211> 366
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30636

ttgcttcctc cagaagcaaa agccttcttg aggaatcttc tanacggccc aagtgggctt 60
ggatgctatt tgcaccccca ttnttactaa ttacaccccc ttgctttttt tggtgattct 120
ttattcgtaa agttatggaa acttacgaaa tctcgaacga tacttggtat cttttcgtaa 180
tgttacggaa ccttgtggat tacataatca tccccttatt gacttaccga atgtatcgga 240
acctcactaa ttgagcaacg atgcttccat ttgaatatcc gcgtgtcacg gaaccttgct 300
gattgtgcat caatattttc tattgatttc tggcacgtnc cggaatttca caaattgccc 360
aatgat 366

<210> 30637
<211> 120
<212> DNA
<213> Glycine max

<400> 30637

atgaatgatg tgataatatg ctgcactaaa aatcttaact cgtaattgac taactacaaa 60
atgcatatat tatatcatag actacactcg atattcttga atcatctaata ttgagtacaa 120

<210> 30638
<211> 426
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30638

cgcatgttat cnttactaat agctatagac atatttgaag accaactata ataatgaatc 60

tggaaattcc tttcccaacc cctcagccag gtccatatga gaaacagagt gctatctgtc 120
 agtctagtga tatcaaaggt ttgattctgg aaaattatat catttctgag cctccatatt 180
 gaggttgtaa ctgctatcca ccacattgtc cttcttatgt tagcatcctt tgaaactgct 240
 gaggagaaat gctgaagaca attgtccaaa ggtctgcagt gaaacacctt ttcttccttt 300
 atccaagata ggaattccca ccatataggc ataattntgc cacaagtga gagtacgtgg 360
 gaagcagttc aggttgacta tgacagaatg gcataaatat tatgcacctg aatctgcctc 420
 ttaatc 426

<210> 30639
 <211> 349
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30639

gcttagctac acanccccct ataatagcta agctcaccn catgatataa tacatgaaat 60
 acttaaaagt ccctactaca aagactactc aaaatgcctc gaaatacaag gctataaccc 120
 tatactacta gaatggccaa aatacaaggc ccaaacaag gaaaaaccta ttctaantat 180
 tacaagata agcgggctca aacttagccc atgggctcgg aatctaccct tangctcatg 240
 agaaccctaa ggccttcctt tggatctctg gcccaatcta cttggtgtct attatccaat 300
 gcccttggtg ngtaagatng catcattccc tccaccttgg aaaggattt 349

<210> 30640
 <211> 313
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30640

agcttgtag aactatcatc acatgacgct ntattggcac agaataagtt gctttctaag 60
 caacttgaga ttttaacaga aacacttggt aagttgccaa ctaaactgtc tattggtcaa 120
 cttcacatt cttctatctt gcagattaca gggttggtga tcaagtggcc tcagaataat 180
 taagaaaggg gggggggggt gaattaatta ttctaaacc ttactaatt ataaaattac 240
 tcttataagg cttatactat gttgttaagt gaataaagag tagaagagaa acttaaccaa 300

cagttaaagc gga

313

<210> 30641
<211> 382
<212> DNA
<213> Glycine max

<400> 30641

tataaaaccc agcttgggaa acaattagaa gggaggagaa ctgatttggt attttatgaa 60
tattatgtct tataacataa gggggcttgg aggtagactg aagaacaaag aggtacatag 120
tttaatttct aaatataagt taaatgttat ttatgagcac gagacaaaat tggaggagat 180
caatagtagt ttatattctt tgttgaggag ttgagatgat tatgagtttg attctaaaaa 240
atcagagggg cggttatggg attttatgat gtggagaaaa gatcttttgg ttgtaaaaga 300
ggtggtgtat aaggaacatt ggttatgatt aattggtgta tgtggcgttg agcagattga 360
agtgtttatt gctgggggtg at 382

<210> 30642
<211> 377
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30642

agcttgctaa tgtgttgcac accaatgagc tcgcaagacg gctaaagggtg atgacttatt 60
tgagggttact gagtatatgt tatacctgct agtagtctgt ttctgtatgt gttgttctgt 120
tttatttacc cctgcaaaaat aaaggaaaca tgagaacagg gaaacagggg actaatccag 180
cttatcagga aaatggttgt ggagggtggc tactgacacc aacagtttgt gagatgctat 240
ttgtttatgt aaaacaaatt gttgtgagtg gatcaactgg agttgggggtt tgtaagatan 300
tttctgagg gtgttggttg ttgataaaag agtaaccttt gganaattga attcttattc 360
atatactctg aaatgac 377

<210> 30643
<211> 252
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30643

ttatgttgca natatttaca atagacctcc tcaacctcat tatctaaatc aaccacagtt 60
tatcaattat gacctttcca gcaacagata caacctgga tggaggaatc accctaacct 120
cagatgggtcc agcccttatac aacaacaaca gcagcctgct ccttccttac aaaatgctgc 180
tggcccaagc agaccataca ttctccacc aatccaaca cagcaacaac cccagaaaca 240
accaatagtt ga 252

<210> 30644

<211> 380

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30644

agctttcttg ataaaacttc cttgagaagc ttctttgaga aaacttcctt gagaagctag 60
agcttagcta cacacacccc tctaataact aagctcacct ccttggaag cttccttgaa 120
aagattccta aagaagctag agcttagtac acacacctct ctaatagcta agcttacctc 180
cttgagatga gaagctagaa cttagctaca cacccttat aatagctaag ctcaccctta 240
tgacaaaata catgaaaata caaaaaanag tccctactac aaagacaact canaatgcct 300
cgaaatacaa ggttaanacc ctatactact agaatggcca aaatacaagg cctaaacgaa 360
ggaaaaaacc tattctaata 380

<210> 30645

<211> 346

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30645

tctcccccaa ttntctataa atagggggag aagtgatgtt gataagggtc cagcccctta 60
ggcacttgct tctctttcga atttgctcgg aaaaattgtt tccgtgaaga aaatctaagc 120
cgaggcgctt gcgaaacgtt tccgtatcgt ttccgtgag gaatctcgca aagggttcaa 180
ccgtttctcg acgtttctca ttcgttcttc atcgttcttc gatcttcaac gggtaagtac 240
ctcgaaccaa gcttttctat tcattctatg taccgtagt ggtccacatt gtgtttcgtg 300

catttatatt ctcgttttgt tactttttat taccctgtt gcatgc

346

<210> 30646
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30646

agccttgaat ttatcttttc tatgtcttga attgaacctt atctaattat caaatattat 60
aatcacatt tcgttgtggg ggaatgtgag gctcanagaa cattgtgtat tttccatctc 120
tctttgcagt gcatatgata gagcggtat taaattccga ggagtggagg cggacattaa 180
cttcaacatt gaagattatg aagatgactt gaagcangtg atcaatttgt gaatatttat 240
attttgtnt atcttatctt gaacagtcac acctcatag tataggatca ccttatctcc 300
tacagttagt gctatntttt ctgtcttgaa gtactctcat gaatttgta aatgcaatgt 360
taatagatga gcaatcttac 380

<210> 30647
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30647

tctctactac caagagcagc tcaatctgtg atataaaata atgtaagaga atcaattaac 60
taacacccaa taagaacatc agaggaagtg atgggggata aaaataaaat gtcatgattc 120
atatacagac agacaaagat gaaagtcag cataggctta ttttctgaga ttntgtgagg 180
canganaaaa gacaggcaac aaaccctttg acagactcaa aatgacaaaa tagaaagaca 240
acaagacaaa agggcaagta atttgatggg gaggagctaa ggaacatact gtgttgaaaa 300
ataaaggggtg aaattaaatg atntaggtag ttaataagt aaacctgagc angagaaata 360
aatcataat atatgcaata ttgcatggaa t 391

<210> 30648
<211> 348
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30648

tatatgccta gatatcaggt taatttaatt tagcttatnt tcttattang ttttaattac 60
tttttttttc tgtcacaaat tactatatat gcactgtatt tcatttctta ggccgcatta 120
attagtatgc atttgtttaa ttaatttcta aatatttcca cttctttttc atgtgtgcta 180
ctacacagtc tacactatag tttgtgtatg tatgagaaag atgaacacta ggttatgacg 240
tgtacgagga gtttgaactg tgagtcanat tactctacaa taatagtaat atctaatttt 300
atcctctata acaacgcaaa gctagaatat accccaatt acatggat 348

<210> 30649
<211> 379
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30649

gctattattg tgtgtagagg gagtgatgaa cataggatgt cttctctcac aacttctact 60
tggttctttt gggatcttag gttgaaagac attgattagt gaatgttttg agaaatgata 120
cacctctgta ttttgacaga gtaactaccc acgagtttga cagctaatag atctcattac 180
atcattcaca tatgtcattg ttatctaaca ttgttccctc agtctatggg ttacaacaat 240
tgtatattct tctctttatt ctcatatagg taccatttga agaagctcca gagcttggtg 300
atggacggta tgtgtttatt aatcaatgat atgcatatgt tgcaatgaat canggtcact 360
agtgtcatta cttctattt 379

<210> 30650
<211> 247
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30650

gctttcactt tccaagcatt aatttccaac atcatgaact acctaactag gaaacagagt 60
aaggtnaaaa atctgccaaa acacattcac atcttacagt ttcttactca aataccccag 120
tacattcctt tgtccgattc gtaccgtgga tnacttgaaa tttactggag attctagtca 180
taagtnacat tntgaccgtg ggatctgtag aaatgtcaga atcaatatgt actacctttc 240

cataacc

247

<210> 30651
<211> 422
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30651

gcttttcttt gcttggctac aaaatcattg gtttgtctag gaacaaagga tagatcctaa 60
gagagcacia atcctatact tatccaagt atcctttttt atatacaatt gcttactcac 120
tagcttttca ctttcatttg cttttgacct tattgcatta gcacacattt cttttgattg 180
gtttctttat tttggttctt cttctctatt ttttaaccaca caacttatgt gttggggagt 240
ctgatgctat atctatttct ttgcatccca attagtttca cctcccaaaa tttggggtaa 300
atttgccttg aaccatatgc tctcctacaa tctaaacaag gtatcttggga gataatcatt 360
taggttcacg gttcaattat ggacaaaatc attcagctca canaggggtgc atatgataca 420
at 422

<210> 30652
<211> 423
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30652

agaccncgtg ctgttaaaat tgaganccat gnagnnatct tgacacttta gtannaacnn 60
naannnnngg aaggagatg tatcgtaaac tgaagcgaat ttctgttact tcagatagtt 120
ttctttgtga gtgtgctaga ataggagctt tgacatgtgg aaaatagatc catgctcatg 180
cattgagaac tggatatct ttatggttta taccatgcat atacatacgt tgttaggggg 240
aaatagaata gctggaaaaa tttttagtg acatgaataa acatgaattt tgtacagata 300
tgaaagtgga acgaccatgc tactggcttt ttaagatggg agtaacgtga actatgaatt 360
cttattcatg ttggctgcgc acaagttggg caatgttaaa attttcagat ggtagactt 420
ttt 423

<210> 30653
 <211> 312
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30653

atacttgtaa tttgttgaag ttagtgacac ttcgtgaatt atcgataact agacgtagtc 60
 tcgggtggttg agatacacta gtataaattt cttgtgtctt attctctctt ctattattng 120
 aactggccta cggtttgaat gtgatcttcg cttttgaaca actctatttg cttacaaaga 180
 tatgagacta ttgtctgac tgtcttgcaa gaattgatat ctatgttctt angtgtactt 240
 catcaacact atcttgatgt attcaaaaag gtttgagttt ataaatttgt aatgttacat 300
 acatgatttg at 312

<210> 30654
 <211> 324
 <212> DNA
 <213> Glycine max

<400> 30654

tcaatctgcc atagctactc agttcgccaa tctccctcaa atgctactac acaacctaca 60
 acaagctcag caaaacatgt tgggtgttgt catccaacca cctcccatta tccaacaaca 120
 accaactcca agtatattgc ttgcacctgt tgaaggaaaa ccatcatcac ccacaataac 180
 accaccagtt tcagcaccaa caccaccacc gaccaatcaa gagtgacgat gatgtcccat 240
 cataattact ttctgtcaat gacaaaagga agagaatgtt atagaaattt gatttatgta 300
 aatacgacac tcttataaaa taca 324

<210> 30655
 <211> 117
 <212> DNA
 <213> Glycine max

<400> 30655

ctgcaacttt acttggtttg ctccggactt aaaccctgtc gacacactaa tttaaaccct 60
 cccctcttta cagcaccctt ctcatcctaa actactatcc ctggcaagac gactagg 117

<210> 30656

<211> 338
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30656

agtccatagt tccaatcaat catgctcagt atgatgcatg cacctgacct caactctcaa 60
 acgtaatgtg gtaccatccc caaggaaata gtctaagcgt tgtagaagca aagcttccaa 120
 gattattttg atgatgccaa agatttttaa aagatgcatt caaacaagat taaagaaatc 180
 aagaagattc aagtgaagat tcaagagaag actcaagata tgcaagaacc tcaagaatag 240
 ctcaagatga gataagaata atntttcaaa gaaaagaatg atagcacaat ttgccaaaga 300
 aaaatcttnt accaaagttt ttactatct ggtaatcg 338

<210> 30657
 <211> 326
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30657

aaaaatttct tggatagatt gaagtctttg gtcataagtt ttatttactt atgctgccac 60
 ttgtgacaac attacatcat acccaaactct gtgtgcaatt tcgatgtttg cagtgtggat 120
 ctgtctatac ttctctcttc aattcttttc aaatttcata ctcttcccaa tagagttgtt 180
 tgcagtttca ctagcagacg tccaaactaa taactatcca ataaatttca ccatcttata 240
 ttcattgattg caaatctatc taccggctaa gataaattat aatatagggtt ttctaagtga 300
 tgtgatctct ntgactaata tgacag 326

<210> 30658
 <211> 434
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30658

agcttttact tatccaagca attcaatttc caaacatcat gaactaccct aaactaggaa 60
 aacagagtaa aggtagaaaa atctgcccac aacacattca catcttacag ctttccttac 120
 tcaaataccc cagtaacatt ctctttgttc cgattcggtt accggttgat cgacttgaaa 180

atcttactgg agattcctag tacataagtc tacattttga ccgttgggat ctgctagaaa 240
 atgtccagaa tccaatatgt actacctttc ccataaccag caatgcacaa gcatttttct 300
 gcacatttgg tcaagttggc tgcacaattt gacagctttt tgctgcacaa tttggcagat 360
 ttcgaaattc ctcttacctc cantccaatt tgctcanatt ggantcctac agtcctaaat 420
 catgcataaa tcat 434

<210> 30659
 <211> 317
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30659

gctncttata tggagcacac gagaccagac aaacatgtca cttcatctat caaagtacat 60
 attgatctac atcatgcaaa tttatgtgac acttctcatg cctaattgaa tgatntgatc 120
 gatctatcaa cgctctattt atacatacat aaaataacaa gacgatttaa ttcctttgac 180
 acggttctgt ccatgatgta caacaagggtg gttacataca cattatttaa ccaattaatg 240
 aaaataataa tatgtttcat caaaactgac ctgatccttc tcatgcttat tgagtgaagt 300
 gattaccggg cttgact 317

<210> 30660
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30660

agcttgatg tttgcttcta tnttctccca gaaaagtata aatataagaa acataaacia 60
 gaagctctct tgttttggct ggccattttg acatattcgt gcatatttac atagagacta 120
 acccattgct aatagtctat attgagacgg gtcaatgggg gtttatataa ctatgttaat 180
 tgctaatagt caatgcctat cagtatcatc acataatcca atgacctag acttcattgt 240
 ataatagtaa cacaatcata ttaacataat aatttacata atatgggtgt cattatgagg 300
 atcaatctct cagacaanaa gtcaaaggaa ggcgggacac aaggacagat gcaatntaaa 360
 catccaattt gttttatata tttcaatgag aaagagatga tatatcatca ctttgacgtt 420

caatgtatga caa

433

<210> 30661
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30661

ntaaaggatg ttntatcagt acaaaaatat atgtntttgc tctggtaatt gattaccaa 60
tattgtaatc aattactaga gatacattac cagagacaaa ttacataaag gctttttcaa 120
aaagaagttt ctcttttgaa atttgaattt taaatgctgt aatcgattac cacttgtatg 180
taatcgatta cctgtgatga aatttcagaa gttaacattg aaaagtcgtg acctttcaa 240
acataactat gtaattgatt accaagaagc tgtaatcaat taccagtgaag agaatttttg 300
aaaaatattc tgaaaagtca cgtgtctntc aaaagttttg aaaagccacc aaggacctat 360
aaatacgtga cttgtctacg aanaacatta gagttnttca ttagaaccta ngtgacatat 420
tctctcaaaa caaatcatt 439

<210> 30662
<211> 411
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30662

gcttcttgtc tcanaaatgg cctttgcaac ataatgcgc attgctctta tggccttcag 60
ataatgaggt atgatataac tcagaancaa atgaaatgaa gatcttctgt cttaattgct 120
taaaaatttg ttacgttgag acttaattaa ctattctctg tttatgggcc atttatttta 180
ttttgacct taggataatg cctccgattt cataagggtta ggtggaattc agagactgca 240
ggatggagaa tttattgttc tagtaagttt tttctttaat gcacactnta actnttatta 300
cagggtttct gctcttacc ttggatgcct aaattaaaga aacatctgaa catttgtaat 360
ttanagaacc gngngaaaat agattcatct ttgacctata ttgacctta t 411

<210> 30663
<211> 387

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30663

 ctatagaagt aaacaagata taaatatacct cacattatat ttttagcatat gtgcataaat 60
 aacaaataag tcataagtca tcaagacata aagcatttgt ctgaggccct ggcattctaca 120
 agtcctaatt ctcttctaata ggcgtagaaa gagccttttg ttagtggttc tgtgaagatg 180
 tctgcaacgc tggtaattag tatctacaga tgctcaaaac acagtcacct ttgtcctcag 240
 actaacggct aatngaccat caacattttac caagataagt ttttattgac atagaagggc 300
 ttatcatatc aggatacttt atttgaaata catataataa ttttgaaaag cataaaaatt 360
 tttatcaggc catcaagtat tcgacat 387

<210> 30664
 <211> 333
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30664

 tgctgcattt cggaatacca ttnttcgggg aaaacatgat gtttgacaga gaaaagaata 60
 aatacaatac ggcgaaagta atgttagaga cagacaaaag gagttgcaag acatcattat 120
 tattgtgcat aataacaagt tttctttttg gtgcagtgca taataacaag ttagtaataa 180
 tcactacatg ttttcttttt cagttgtcgc ctttattcat cgaagtatga ctctaattctt 240
 gagtcttttt ttttggtata aactaatctt gaatctgaat ggggtggttaa gtaaatttct 300
 aattgaaatg atactttaat ctaaaattta aac 333

<210> 30665
 <211> 426
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30665

 agcttaacat gtagatattg gaacacatca tcgaaagcat cccacaaggt atgtgttcag 60
 ttctgtaagt taacgagggt gaagtataac ttctcaatgg ttcttttgaa aaattgcatt 120

tgcaggtgca agaaagataa ggactaccta tataagcatg aagtntaacc gcttcaagaa 180
 gctgggactt ggctttgata tgcttatgaa ggatagggac acaggctagt aaactagtgt 240
 ccaacaagag taatgttata aactattgtg cagattatct tcatgtattc attatgaata 300
 gaaagggttc aatccttagt gacaccctga tattcgaata tctgaaacgt gtaatttggt 360
 aagatgaaat tcaatcgtca cgatatttca tctatgcagt aatttgtggt atgttatgac 420
 tttggt 426

<210> 30666
 <211> 212
 <212> DNA
 <213> Glycine max

<400> 30666

actgcatgta ctgctgctct aatctgactg tatgcatttt cccaaaggat aatttatgct 60
 ccaatataat acgaaaataa atgtcttgaa agaattgaaa atgtattata gaggatctca 120
 atccaatgag atactaattg ataagcctat tttaacctct acctaaaata aaatatacaa 180
 gatctaattct atatggctta attcgatata ag 212

<210> 30667
 <211> 182
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30667

cacattattt ccatgacaca tatgcaaaga tgatgatttg ganatnttat gcanaactgg 60
 tcatgcatgc acctatgcgg aactcaagt gtcaaatttt tatggatcatg ggatgctacg 120
 gctcangatt catttcctct attggttagtc aaccaatgt atcaaaatat gttcttttat 180
 ca 182

<210> 30668
 <211> 348
 <212> DNA
 <213> Glycine max

<400> 30668

tcaagctagg gccagattct cgtgcatgca gaggcttctt ctataaaaac tccaaactcc 60

ctttgcaa at ctgatttcag gcttaa atag gtggccttgt t cgtgctcgt gcgcttagcg 120
cagatctaga t cacttagcg cgccta agtg gattgtggct taacgtgctt gtttcgctta 180
gcaa atgagc tgaagcgggtg cacttgatga cctggagtgt gacacccctct accccgacat 240
atatataaat aaataaaata tataaaaata tattggtaaa caaaatcaca tgggtaaaag 300
gttcacattc acttcattta ccaaataaaa cttattaataa acaaattc 348

<210> 30669
<211> 276
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30669

ttacctcctt gagataagaa gctagagctt agctacacac accccactaa tagttaagct 60
cacctccatg ccaaaatata tgaaaatata aaaaagtctc tactagaaag actactcaaa 120
atgcccttaa atacaaggct aaaaccctat actactagaa tggccaaaat acaaggccca 180
gaagaaggan aacctattct aatatttaca aagacaagtg gacccaacct tgacccatgg 240
gctcaaaaat ctacctgag gttcatgaga atccta 276

<210> 30670
<211> 398
<212> DNA
<213> Glycine max

<400> 30670

ctggatcctg gatcctggaa atcaaatttc ttcttgaacc ttgaagtgtt cttaatggaa 60
tcttgaactc attctttgat tottgagatc atcatctttg gtatcatgaa ttggtgatga 120
tctttgagtt tttttgtatc acctttgtca tcatcaaaac ctctttgaat caatcctgat 180
tcaatatgaa gctggcttct acaatctccc ccattttgat gatgaccact ttctaaatca 240
agaaacacac acacacacac aactcacac actttttcta gccgatgact cacataaaat 300
tcctttctcc ccctttgggtt tttgaatata tgcttggctt aaaattaaag tgattactca 360
tgtgagtcct tggattaatc cctattctct cccctttg 398

<210> 30671

<211> 303
 <212> DNA
 <213> Glycine max

<400> 30671

gcattcgcta agcacgacac tcctgtgcta agcgcgagga agaatccaga agaagatgag 60
 ctgtacaagt tcactaagcg caccacttca gttcatccac taagcgagaa aggcgcacta 120
 agccaaaaat cactaacgtg cgctaagcgg tccatacgtg cgctaagtgc acgagcacga 180
 acaaggccac ctatttaagc ctgaaatcag attttgtgaa gggagtttgg attgggattc 240
 agagctttgc atgtctaggg tttctagaga gagaaaggtc caagttctag agagttttga 300
 gag 303

<210> 30672
 <211> 353
 <212> DNA
 <213> Glycine max

<400> 30672

agctttcact tacatacgaa gatttgtgag tgacaaacca tgttgtcatt cttgacaaga 60
 taaccaagag gcatgtccat gtatactccc tcaatcaaat cactattgaa aaacacatta 120
 tttaaatcaa gctgaaacat gttccaatth ctgtgaggtg caatggaaag aaacactctc 180
 attgccgtat gcttggcaac aagtgaagaa gtgtccaaaa aatcgatctc tgcttgtatg 240
 ttgttgtgtg taaccttttg caacaagacg agccttgtat ctatcaatgg agccatctgc 300
 tctatacttg accatataaa tccatctgca actgatgggt ctattatcgg gtg 353

<210> 30673
 <211> 229
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30673

tgcattatca acttctcatc cctggaatat gcacccatct ttcgatagta cattgttagg 60
 tgattcttat ggcaagtcgt ccccttgtac ctatcgaaat caggtagctt gaactttgga 120
 gggatgacga cgtccggcac caatcaaagg tcggtcatgt ccgcaaattg ataatcgctg 180
 aatacttoga cagccctcaa cctcttttgc atgagatcga gntttccct 229

<210> 30674
 <211> 364
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30674

agcttttcga ttcattctat gtaccctggg tggccacat tgtgttttat gtatttctat 60
 tctcgcttca tttacctttt ataccnctc ttgatgtgct taagccattt tacttaagtc 120
 atttctcgct taacctaaaa ataaaatata atttcaccga tcgcttgaat tgtattatcc 180
 gtttaacttcn gttaaaatga attccgaccg ttcggtcgtg ccgtaaccac gttggatata 240
 ataaatgagg tcacaaataa tataataatc aaaaaacatc tctttagtaa ataaagcgga 300
 tatcaatcgg acgtttctct ttgggattct cattcttatt gaatngctaa taactaagtg 360
 aact 364

<210> 30675
 <211> 186
 <212> DNA
 <213> Glycine max

<400> 30675

ttagtgaaaa catgattaca tatctaggat ttttttgctt gaatgttggg aataaggggg 60
 gttttgtcat tggatacagt gtgatggctg cttatgatta tttgaccatc ttgagtcatt 120
 gctatggtaa atgtgacatg ctgaatatag ctgttctaaa gctacatgct aaaaatcaaa 180
 aaaaaa 186

<210> 30676
 <211> 370
 <212> DNA
 <213> Glycine max

<400> 30676

gtgtaatcaa ttatgatcag attgtaaccg atgaaaatag agtttttaaac attgaagaaa 60
 ttttctaact ttagaacctt tcttcttagt cctacatgat gatgcatgat gcacgtatga 120
 aatgatagag actaagatgc aacacacaat acaacagtca atacaaacgc cactcaagag 180

agttgggcat gtaaaagaca aaacttcttc aagttcttct ttaagcttca aggccaagtc 240
 tttattttgc tccccttata tctaacaatc tcccctttt ttggctttga tgatgccaaa 300
 cttgaattct ccatttgagt gcatttggag agtcttaaga gtagagactt ttcttagaca 360
 aacctgaatg 370

<210> 30677
 <211> 402
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30677

agcttccagt ttattttcat attntcagct cctagtttat agcttataag ctttcaacta 60
 acttattaat tagttttacc aaattaaatt tgtagttta taagttttac cttagttata 120
 aatgaaaaaa taaattaagc taaaataaaa tgctcgtctt ttatgtattt tttgtttcta 180
 ctctgctcct ctaatttagn ctcttataac tttcggaag ataatagaaa atggaatcga 240
 ttgaaacata gtagaatggg tgaatcatga atagaaagaa ttaacaatat gtcactctat 300
 tattaataaa tgtagatgta taatataatg gtcaaaaatt agattccatc atttgataaa 360
 aggagttgaa taaattatct ttttattcat tatgaaaaat aa 402

<210> 30678
 <211> 269
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30678

cttatttcat gtattcactg ttcctctggt acttgattat tatatatatt tatctttgcc 60
 gagcaaaaaa caaatgtcta tgggcctaga gcatggcaat gcaggtgacc canaaatgga 120
 tctaaaatag actctgaaat cattntagaa tttgggctta gtgaaaaggc ctaactcatc 180
 ccatataacc gacttgtagg gtgaggattg ctcaaacttt ataagctcta tttaagttat 240
 atctctagac tatgtgggac taaatactc 269

<210> 30679
 <211> 417
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30679

tgctttctct tgccatttcc tgtgaaggca aaaatttgga aagttagttt taccagtggg 60
acactactct taaaacaaga atggcataca acctcctccc ataaatacaa acatcaatgt 120
aaatttagag caagcttatg cgcattgttc cttacgaacg ttcacttgcg gaagacatcc 180
tattaactaa gaaaaatgca cccatataca atcaaggtag cttcattacc tagattattt 240
acatgtacct ccaagggtga tttgttattt acatcacaca cacctccttg gctgaattta 300
catacatgca tactcaaagc attntggggg accaaaaact gcacatggcg tcactcttgg 360
atntctaaat accctacata tacaaaacttc acgatgaatc ttgactgcct acacaat 417

<210> 30680

<211> 421

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30680

agcttcaatt ctaccactaa atgggtgtaat aagttcatat ctcaactntc atacctctac 60
attcattttt ctttctaate ctctatcaga tcagcactcc tcacatcttt caccttgaat 120
tgacaaaacg tgacctatat ttctatgtgt gtcttctgag gtatcattcc ctaattaatt 180
ttacattntg acaccttttc ctctctctct ttcaggatg ctgctagtta ccgtgatgag 240
ctaaacaata ttgccccaca ctctctttta aaatgttgca gcgatgctac aacattggta 300
tgatccctac tttcagttaa atacgtttta tctccgagtg atgtaagcta tctcaagaga 360
aatacagtta acaaggaaaa ccaacttcct tttttaaaca gtcttttaag ttgattgcac 420
t 421

<210> 30681

<211> 168

<212> DNA

<213> Glycine max

<400> 30681

agatattcca aactatttgc cctaattgaa aatctatttc actttgtact caagttatga 60

attaccttaa tgacgatctt cttaagtaaa tgaacaatg tggatatgaa tataaagcaa 120
 ttatgatata aggagattaa gggaagagaa aatgccaaact cagtttta 168

<210> 30682
 <211> 363
 <212> DNA
 <213> Glycine max

<400> 30682

ttgcttgta ttgtgagaca tcagaggcta gtatttgaat aaatgtgggt aagaaaaatt 60
 caccaaattg atagagaaca atctaaaatc atacatctta ggcaaataag gcatgctagc 120
 cccaacatt attgcatttt gattccatct ttacacattc aaattgttgt ttatttctcc 180
 tggtatcttt tctttgcct tagtctaaat ttcaaactta caattccgta tctctttctt 240
 cttttgtttc tctcatttc ttaataattg gatttgcac acttaagtac aaccaagtc 300
 cctctggatt taattgttga acttcaattt caatctttac tactcgtgat aaaattacga 360
 cac 363

<210> 30683
 <211> 290
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30683

cgaccttaga atactcagct tctcgctcan aattcacttc ttggttggtg tttttggttt 60
 gtgctaaagg tgggtgtcgt cattggaagt gtggtagaca gactttgtgg tagatttagg 120
 gatggccttt gtggataact ggggtgggtgg taaggaggag gtttggttatt ggctgagtaa 180
 tgacattgtt ggggttggtgg gaaacttggc cgtataggaa tggcagtcac agcatgggtt 240
 tctccctctt tctcaccctc ttcatttggc ccagttttct cagtcgtcta 290

<210> 30684
 <211> 498
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30684

agacaacggg ggaaatgagc gactcacgnc gtncnaanac ncngnacccg ggatcctgtc 60
 agtcacctgc ggcatacaag cttcagttta ttttgcattt ggctgaaatt aatttaagtg 120
 ttgtattgcy tattccatga tgatacatte tgtgtgcytg ctaaataaat tggctcactg 180
 cgatggctta tgagatgggt tgccttcaga aaatgggtgat tgcttatatt atcttagaga 240
 ttgctgatga gaaatgggtg accccctgat aacacgcata tgctgtgatc gctctcgtgt 300
 gcttgctata tgcacccatc accatatcta tacatcctat gactgcttta tgcactagaa 360
 ctggtcataa gactttctga gaatatatat cgttatgcat cgggcgttta cacacngcc 420
 cggctttaca gaacatcatc ccaaaaaact gacatggctc tcttggtgaa tttccatact 480
 tagagtttga ctctctcg 498

<210> 30685
 <211> 338
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30685

tagaatacta agctntgatc catgaanacg acaattatct ttttctcaga tgtcttattg 60
 agtccaataa tataacgaga cgctcgaaat tgantgttga agctctgagc taattcaaac 120
 gacaataact gtttactcgg atgtctgatt gagtcccgcc atatctgag acgctcgaaa 180
 ttgaatgggg aaactctgag ccaattcaca cgacaataac attttatggg atgtgtaatt 240
 gcgtcccgta tcatatcgag acgctcgaaa ttgaatgggtg aaactttaga caattaaacg 300
 acataacttt tacttgatct ctgangagtc ccgaacat 338

<210> 30686
 <211> 419
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30686

tagctttctc tgggccattt cctgtgaagg caaaaatttg gaaagttagt tttaccagtg 60
 ggacactact cttaaaacaa gaatggcata caacctctc ccataaatac aaacatcaat 120
 gtaaatttag agcaagctta tgcgcatgtt tccttacgaa cgttcacttg cggaagacat 180

cctattaact aagaaaaatg caccatata caatcaaggt agcttcatta cctagattat 240
 ttacatgtac ctccaaggtg tatttggtat ttacatcaca cacacctct tggctgaatt 300
 tacatacatg cataactcaaa gcattntggg gtaccaaaaa ctgcacatgc gctcatcttg 360
 gtatntctaa tacccttaca tatacaaact tcacgatgaa tcttgactgc ctacacaat 419

<210> 30687
 <211> 328
 <212> DNA
 <213> Glycine max

<400> 30687

ccttgccctca tagaggtcca ggatggactt tgcagccgaa tgatctagtt ccgctccgga 60
 gtatgacagt caccgcttta tgagcgctgt acaccagcat cgcttcgagg ccatcaaggg 120
 atggctggtt ctccgggagc gacgcgttcc tctcatggac aacgagtatg ctgatttcaa 180
 tacgaaatag ggcgccggcg gtgggcatca ctggatactc ccatacccag tttgaccgca 240
 aatagtcctt gagtttatgc catgcttggc cacaagatg gcgtgctgac atagatcctg 300
 cgtagggctca gtgaatctgt ttgtgcca 328

<210> 30688
 <211> 349
 <212> DNA
 <213> Glycine max

<400> 30688

acttgatgat tggtactgtt tctcaattat aagataagca ttgtgttatg acttggttct 60
 gattatcaat ttaagataat gatgactcct tcataactct ccataactga agtggtatgt 120
 aaaggtgata atatgtaatt tgatagttat ttaaggaaaa aatgcctaag tctatactaa 180
 aataagttgt cattattaaa atgatgttac aaatccactg attatatttg ggggtgaggt 240
 caatggctca agtatcacat tgaccactgc aagattttac ttcattgttaa tatactttaa 300
 ttctttatct gatgcaattt aatctatcta tcgcttatga ctacttaaa 349

<210> 30689
 <211> 273
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30689

ntgcggatgt ggtcttcgcc ggtgaattgg tcgaagcgga tttgaaaaga ggaaaatgta 60
 atcatcctgc ttggacgaat gagaaaattg gggcaaatga agatggtgag aatgaaagag 120
 aaacccatcc tgcgactgct gtttctacat gggaactccg ccaccagctc aacaatgtca 180
 ttacatagca aataacaacc cttctccgtt actaccacct aattaaccac aaacgccatc 240
 ccttaatcat ccacaaaacc cacctgtcac aca 273

<210> 30690
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30690

agcttgaagt tatatgttaa aataactnng tgtattcttt ttacattctc ttgcaacagc 60
 ctncacttga gnnttatcat tanannagtg ttcagggaaat nntaatccaa acctattaan 120
 nattgatecc aataanaatt gatnaatcga gcanaattgt tatanaaaaa tacctaaaca 180
 cttcaatgcn agttcggttt tgtgattctc atgttcaaaa ttgaactgag tcaaattgga 240
 aacgtaacta aatttatatt agtcttattt tgtcttcttt ntttactgca attcctatat 300
 atttatcttg aaatacaatt taaccttatt tgaacttata ttattatttc tgaataactt 360
 gaagataatn tgcaatntag tctgtgattt agatcaagtt gtgggttttat gaccagaatt 420
 aatagtttga aatgatattt ctgtatatgc attatg 456

<210> 30691
 <211> 465
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30691

ggatgtgatt ggggacctga gactcaaacn tataaaaccc ngaatggaca tccggtgaac 60
 ctttgaaatt ngaatttatt agagcttccg aggttcaatt tcgagtgtca atatatgtga 120
 tgcgccatat atggacattc gagttaaatg ttatgaccct ttaaatatct caagagctta 180
 cttggtaaatt ttcgagcctc taacatatta tgcgccccag tcggacatac gtgtgaagag 240

ctatggccat tgaacatct ggcacagtta tcgatgataa atttcgagct gatcgggtatt 300
 ataatagccc tgaatcggac atccgagtgt aaagatatga ccatttgata ttctcaagct 360
 ctttccgtga tgcatttgta gcctcttaga atataatgcg cccgatatag aatccgtgtg 420
 aaaagttatg atttataaag tctcagaagt ttcgagtaca ttccg 465

<210> 30692
 <211> 358
 <212> DNA
 <213> Glycine max

<400> 30692

agcttgtttg ttataaagac ccaataattc tacctattgt tgtcattcta tttaccatgc 60
 attttatagt ttttagcata aaagtttagt ttaaattctt tttgaaatta tcacttatac 120
 atgttatctc aacaatgctt caattctgaa cttaattcag gctaacatta acctcccata 180
 cttccatggg aaggataatg tagaggctta tttagattgg caaatgaagg ttgagcaatg 240
 aatgttcctt tagctaccct tagcttccaa gggtagctc tctatagggtg gacttcactt 300
 gttatggaaa gaaacattca ttgggatcct ctaatagagt attggaatga cttgaaaa 358

<210> 30693
 <211> 395
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30693

gagaengaga ccaacatgtt agctatcatc gccaaagtacc aagaagagtt aggtctagcc 60
 acgggccacg agcatagaat cgcggatgag tatgctcaag tatatgcgga aaaagaggct 120
 agaggaaggg tgatcgactc ttacaccaa gaggcaacca tatggatgga cgggttngct 180
 cttacctga acgggagtca agaactatca cgcttgtag ccaaggccaa ggcgatggca 240
 gacacctact ccaccncga agagattcat gggcttctcg gctattgcag catatgataa 300
 cttaatggcc acataataga aatcgtatgg acttgatggc tctcaacctc actgatacga 360
 ctctttttga ataaatgagt ggtcatgttc tctcg 395

<210> 30694

<211> 207
 <212> DNA
 <213> Glycine max

<400> 30694

aaaattgatt ctcacacccg aagctgtgcc tttatggaga atcctccttc ggcttatcga 60
 ttctatgtgg ataatggcga cagactgtgc atattcttca tcttatgcat attttctatt 120
 gttctgccct tgagctctca gaaagtcaac aatgggtgggt cttgaatttg catcctgcat 180
 gatacatacc aagtgtccat ggcttgg 207

<210> 30695
 <211> 341
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30695

agcttgtaat tgatgcttaa tggaggaaaa gaaagaggga gacaaagaga gatggcgagg 60
 gcacnaaatt gaaggaacaa aagagggaga gaagtggaac tttgaagtat gtctcacaag 120
 actctcattc atcaaagtta caacaagtgt tacacatgct tctatgtata gactaggtag 180
 cttccttgag aagctntctt aagaaaactt ccttgagaag cttctttgag aaaacttgct 240
 tgagaagcta gagcttagct acacacaccc atctaaaaac taagctcacc tccttgagaa 300
 gcttccttga gaagctagag cttagctaca cacacccatc t 341

<210> 30696
 <211> 341
 <212> DNA
 <213> Glycine max

<400> 30696

gaataccatt tttcggggaa aaaatgaaat cgacagagaa aagaaaaaat acaatacggc 60
 gaaggtaatt atgcaagcac aactatggga cattctcttg tgggtactact ctctgttgaa 120
 atagggattt cacataactg atattgaatt tggctcattt tttatagacg atctgacatg 180
 aactctcatc ctttggtgct cctctaatec atcgaagtat gactctaate ttgagtcttt 240
 ctttctggta taaactaatc ttgagtctga atgggtgggt aagtaaattt ctaattgaaa 300
 tgatactcta atctaaaatt taaactccaa aggggtataac t 341

<210> 30697
 <211> 343
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30697

 agcttgtata ttttcccaat tcatggntat tttggagtaa attttgtaaa taaatcttgt 60
 tttatggtta acgctgtctc tagaagattt ccattggatt taatgatgaa atctgtgcat 120
 tctcaagtga aaaaaaaggc taagttttga attgcaaaaa gtagcagttg ggctaagctc 180
 aacagttggg ctaagcgcaa cttcagcgcg cttagcgcaa aggagaattt ggtagagcat 240
 cagcatcaaa gttgcgcgct aagcgcgaga ttagtgcgct aagcgtagta ggtgccttca 300
 gccaggctaa gcgcgaaact ggtgctaagc tcaattccac tta 343

<210> 30698
 <211> 208
 <212> DNA
 <213> Glycine max

 <400> 30698

 agcttctata ctacccatt tctctccgc tttgggacat cgataagcca aagttcgtgg 60
 caaccaacac aagatgatat aactaaagtg tacataatca atcataagtc acaaccagat 120
 ataagccaat cgtccataag atgaaaccaa atatagtcca agcataaata acgtataacc 180
 aagtataatg caagcgtaaa agactaag 208

<210> 30699
 <211> 401
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30699

 catcaagctt gtgttcgctt agactacatc gcattctacac cttttgtacc aggggcaagc 60
 gagcttghta cagcgagaga ctacatcgtc ttctgcacct tttgtcatcc agagacggcg 120
 agtccgatga catgcggagg taccttatgg ttatccgcac cttttgtcag ccagaggcaa 180
 gcgagcccgt tgacacgcag agactaacat cgctcatctgc accttttgtc aaccaggggc 240

aagcaagctt gttgacacgc agagactaac gttgtcttct gcaccttttg tcatccagag 300
 acggcgagtc tgatgacatg cgagggtacc ttatggttat ccgcaccttt tgtcatccag 360
 agacngcgtg tccgatgaca ttcnngnggta ccatatgggt a 401

<210> 30700
 <211> 522
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30700

tgatattatt tatctanggc atntcccatt ttaccttang tggcggaaga aagtcnaga 60
 gagctgtnga cntctcnnga ttcttattcg cttagaacncn natatcgcat ggggggagng 120
 gcgttgctat ggccctggaa taatcgaaaa catagtatgt agtatgttgc ctcggtanga 180
 aaactaaacc ttgtgcccان agatcccgtc tctctatact tctcattcac cttatgttat 240
 ttcatatcgc agaaaacact cttggctttc catacgcgcg tgctttgtga atgcaaactt 300
 gatatgaagt taccgcacta ctnatcatct tgagcggttat actcaacgaa ccacttgtgt 360
 gaaactggga tgttataata aggccatgat atcgtggagg tctaagtata acgacaactc 420
 gcgaagtcaa tgtgggctta cctgcgaaat acatgtggga catgttacat gagcccaaca 480
 aactcagggg ctcttttgtt tcaactaaga acgaacgtgt tg 522

<210> 30701
 <211> 319
 <212> DNA
 <213> Glycine max

<400> 30701

tcagcaacta tggctattgc tacgcccact ctctctctcc atttcgcaa attccccatt 60
 cgtcaaacgg attcatcttc tccaatacgc catcgcagcc cttctggccg acggttgaat 120
 ctctattacc ctcgtagttc tcaaagggtta cttttttttt gtttgttctt ggttaaatga 180
 aattgaaatt tcgaatttgg attctgagtt aaatgttaac cgggtggttgt ttgatatctc 240
 cttctgttct attggttaca gaattgcgtg ttggcactga tgtgcatttg gatcattgtt 300
 gcaccgatgg ggatgggga 319

<210> 30702
 <211> 506
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30702

aaaaaaacgc ggggacggaa ctgagcanag cactncaann cncnngaac cgggatcctc 60
 tgagtcacct gcggcatgca agctgccatt ggtatttgat ggtgcatca tccactgaga 120
 ctaggcaagg ccggaccctn ctatacaagg aggctatggt attcaaagaa ctgtgtaatt 180
 tttaaaaccc aactaaaatg gtttgagccg tcaaattctg gaacatctag tttcattctt 240
 ggtggatagg ttaagtgggt tcgtggtgca agtatgactg aagacgacta acgtgaagct 300
 gtctgtggtg aagtgtttct atggtggtga cactgtgtta aggtcatcaa gtttctcact 360
 aacgtggaat ctgtagatg ccaatatggc gatagctctc tctatgcgat cctattggat 420
 atggaccgcc ttnattcacg catggtgagt ctctatttga agcacactag atggtctgag 480
 ctgacatcaa tgctatatcc tacgcg 506

<210> 30703
 <211> 93
 <212> DNA
 <213> Glycine max

<400> 30703

gtttgtgggt agtaatttga ctgatatgta ttcaaagtgc ggggagttgt ctgatgcatg 60
 taaagctttt gaggaaatgc cttgttaaga tgc 93

<210> 30704
 <211> 359
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30704

agctntgatc canaatcctg actcaccata naccttgacc caaggtgaga atgccaatcc 60
 ttatcctcgg aagcaaaaaa agaggagaag aaaatttcca atcaaaggaa aaaggagaag 120
 aaaatntcca atcaaagaac aagagaaaga aaatttccaa tcaaaggaaa aaaaggaagc 180

aaagaaattc ccaatcaaag agtgggagaa agaaaaaaag aaaagaaagg aaattcccaa 240
 ccaaagaatg ggagaaagta aaaaagaaga aagctcctga tcgaaagaaa acagaagaaa 300
 tgtgcagaga ggtcttttga cgggacaata tctgaacaat acagaattgt caccaaattg 359

<210> 30705
 <211> 220
 <212> DNA
 <213> Glycine max

<400> 30705
 aacgttctct tgcacaagac atttatatca aagaatgcac ccatatacaa tcaaggcagc 60
 ttcgatcatct agattattta cacgtacctc caatgtgtat ttgtaactta tatcacacac 120
 atctccttgg ctaaattcac atacatgcat actcaaagca tgtaggggta ccaaaaattg 180
 cacatgtgca cctcttttga tatctaatac ctatacatc 220

<210> 30706
 <211> 386
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30706

ctcacattca ctatcttcta catcatattc aaagttgtct aaataaataa taaagtcac 60
 tcgactcata gaaaatcata taagtctcat acaattaata tagaacctat atcctaattg 120
 cacatcctat cagagcgtgg tgttcccggtg tcctctagca tgagggttctt catagtcac 180
 cacctattca tctgtcccc cgaacacaag ttcaagatca tcacangatc caaacacaac 240
 aacacacagg gagtgaagta tcacattcct atgctataga gaaacatgac aattatatat 300
 acatattata taaatgagat accacttgct taaacatagc tcacgtaact tcaccacttc 360
 atcattcaaa attcactctt caatta 386

<210> 30707
 <211> 450
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30707

atttacaaca gtgttacaac agaacctaac tgtntctaata tatatggggcc attaaatcta 60
 tcatgtgttg acagtaattg attagcccggt gaatttcctc tggagctgaa cacacttcgg 120
 ccatggccct tgctgtggct agtacatgcc ggagctcttg acttccattt aagggtcaagg 180
 cgaacctatc catccacatg gtcacttctt gatgcaatgc atcaatcacc ctacctcttg 240
 ctgtcttctc ggcgtatgct tgtgcgaaga cctctactag ctttttctca tgggtcaaag 300
 attggtttaa ctcttctatg tactgcccta atatagctat aacctgcttt gcttcttggc 360
 ttctaagcgt gtagccaaac tattcttggga tctgagcaac cagtaactcc tccttttagac 420
 atgccatgac ctctgattgg tcttttctc 450

<210> 30708
 <211> 430
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30708

tgaaaccttg tacactggaa tccagttagt ccctggatct ctgagcacct gcggagcagg 60
 aattttaact ttgacagtcc agttcagcgt ggcacatttg tatggtgtct gtggcaacga 120
 tgctgtcct gagatagcct acttcatact agatgaacct acatagtgtg gagccactg 180
 cccccggtgg ggctttgaac gctgtacact gaatttagca caatgacccc catcctgaac 240
 tggagcggtg caccgctgg tttaagaatt catgactccc tatgagatat ggcgcgctga 300
 ctattgggga cagtcatgac gttggggaga ttaactgaca aagcgcgctg gggacactct 360
 gatggtacgc gctaactcag tgggttatta caaaggaccg tgtccttaat ggggccgagc 420
 ttggtccgcn 430

<210> 30709
 <211> 221
 <212> DNA
 <213> Glycine max
 <400> 30709

actggccagg cccaaacaca catgcaagac aaagtgagcg aaaccgagca gcgcgcagag 60
 aaaaaacgca gactgacagc tgaaccagta cctgacgcag aacaacacga aatggaaacc 120
 ccagcggaga cacgcacggg cgccccaccc ccgagacag aacagggcga cacaggcccc 180

cacaacgaga ggcaaccacg acccagaacc ggacacgcga c

221

<210> 30710
<211> 299
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30710

agctngcatt atatcttata cgagattcat tatcagatga actatccaag atagcatcaa 60
catacttctt aaaaagggtgc tctccaagga ttttccatct tttggcttca atttaacatg 120
agccctaccc tgaaaagctt caacaagtac ctaccttggt ctaaagagag ttatcttaag 180
aaatctaaca tactcttggg atggaggtgg ngcgctctat acttccatgg ccacattagc 240
aacaacaaca cgtgcctcat catcaaaatt cacctcttca atgatggatt gtacaggat 299

<210> 30711
<211> 247
<212> DNA
<213> Glycine max

<400> 30711

tctatccttc ctaagatgga gcctttccca ctactctca ttaagaacta acggtgtcaa 60
tggattgaac ccatacacia cctcaaaacg tgactgcttg gaggttctat gaaccaccct 120
gttgatgcc aattctacat gacgaacata ctcatcccaa gacttatgga tgcctttcac 180
aagagccctt catacgggtg ataacgacct attcactacc tatgggtgcc catcaatttg 240
tggatga 247

<210> 30712
<211> 357
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30712

agcttctctc ntttcttggg taattattat attctgtttg taagccttgt attttgctat 60
gtttttatga catttgaaca cttagtattt cttttaata tttgtttagt atgactaaac 120
atgatgatta cttgctcttg gttgattatg gttatgagtt ttaaacttaa ttattttgat 180

gatatatgat tagtggtatg tacttttatt tggttattat gaatgactct ctggattata 240
 tgacattcta tgaagtatta tctttctaag atngatgaat gtgtaagtta tcttggttga 300
 tagatctcta ttctcttgta tgattagaaa tttatgtatg tttatatatg tacgcac 357

<210> 30713
 <211> 203
 <212> DNA
 <213> Glycine max

<400> 30713

aaagaatgtg actcttccaa ttgaatatgc atatctatgt tcacacacac tattgatcga 60
 ctaccaaaca gatgtaattg attacatcat ttcgatatta tttggaacgt tgcacattca 120
 gtttgtaagc ttttcgaaaa ccatttagct attggtaatt gattacaata atctggtaat 180
 cgattactag acagtaaata ctc 203

<210> 30714
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30714

tttctacatt tttcaagacc gacggcngc tttgttttag cggacacaat cagacagccg 60
 agngagaagt gatcgccgag cgaacngact ccgacactaa acagtcaata tcgagcgtcc 120
 tgatataatc cgggactcac tcagacatgc gattaataaa gtcgttgctg tttgaatgtg 180
 ctgagatcat taaactttca ttttgaacgt cttcatatat taccgcactc aatatgacat 240
 ccgagtcata agttattgtc gtttcggtct gtaccgaacc tctgcatact gtttcaaaca 300
 tctcgaattt tacgaaactt tttatacatg tgagaaacag tttttaccag tcgtatctgc 360
 ttgcaactct tctattttta atcgcggtta tatatcacn 399

<210> 30715
 <211> 496
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30715

tgaaaccctg ttganccctt gaaanacttg ataccatggg gacacccgca ccatanagtc 60
 acaataccgg cacnagctct ttttcccggt tatttgattc cggccgcaag tttttgttgt 120
 atgctgaggg cgacacgcca cggggcgagc tacttgatgg tatagatcac acacccaggt 180
 cctgtgcatg tgctataaga taccgcacta ctcaatctag cttaatagat gagagcaacc 240
 atggatcaaa aggttctttt cgaagcgagg gatcagatac tagtcgactg gtgacgccta 300
 gccaaagtttt atgcacaaac ttaggacact tgctcaggtg gatacgctcg gtctctcacg 360
 tgcgggatta tcaatgaaaa atgaatgtct ccattgtcgc tagtatatat tgaccgctga 420
 aatgatcggg cgataatatt tggccgtact atagaaggaa atagaagtgt catataaacg 480
 ctcaactagg cctccc 496

<210> 30716
 <211> 453
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30716

tgaaaccttg aacactgaaa acnggtgatt cactggacac gggagactta gagcgactgc 60
 ggcaagcttt ctaattcgtc ggcagcaaca gtgtttctca tcgctctaca tgacatagat 120
 atctacgctg caaacaatg gtactaatac atcatctact cttatatcaa cacaacgcta 180
 ttgggctatg catcacatct ctctgcggag ctctgcgttg gcatgctagc tcattaaaat 240
 gggagcgtag aagcctgaca ccattgctaga gaagtctgga tagcgacgta gttcttttagc 300
 tcttgtagca catcgtgagc tgatgacatg gcaccattgg ctgagagacg cccgatacta 360
 acatattgaa ctctgctgta ctatattaca tgagtgtata tcaagaccaa ggcacggctc 420
 tactgacgga tagagagccc agacactgac tgc 453

<210> 30717
 <211> 348
 <212> DNA
 <213> Glycine max
 <400> 30717

cgccaaccct ggcacttgcg gcaactacgc cgcccagagg ctaactgtga ccttaacacg 60

ctgggacccg cgacagagca gcagttgctc agtatcacac cggccactgc cggcaacgtc 120
 aaacagatga cactagagccc cccggcgacc gacatacagc gggcggcggt caggaaggaa 180
 aagcgcccaa tatgtgcaa ccgaacaccc aactcaatcc gaaaaagggg aagaacccaa 240
 aaccgcaagc cgacgaaata cccgaatcgc cccgatgcga aaacgatcag ccaacgcgca 300
 cccgaacgtc atgatacacg cagaccctc gcacacaaag tcaccaac 348

<210> 30718
 <211> 376
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30718

tgtagattgt tagtgggtgat gaggaaaaag tccctcagca tcgaaggcta gctacatcga 60
 cacgtagaca acaaacaact atagatgttg caaaggatgt tgagaatgtg gataatgttg 120
 ctgatgagcc tcatgaggag cctcacaatc tagttacaaa ggatgtaggt ggtgattcac 180
 agggttttcc aggcggtcc caagatacat caatgttgat gtcatatgtt gatcatgttg 240
 tagccaaagt gtggatagga gaggtagtta tttgtttaat taaaacttat ttaaataact 300
 atttatcatt ntaatttaca tanaataaat ttaattattt ttaaaacaat actgagttga 360
 agttggcctc tcatgg 376

<210> 30719
 <211> 444
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30719

ntatgagaat ccgcattcaa aagcaccatg aatattcaat tatatgattn ttaattatta 60
 attaataata taattaatta actttntaaa cattatTTTT aatttttttt ttcacaaact 120
 aaaagagaga taaaagataa tagaaggctc tgagatgaaa gaaacatata cttttaattg 180
 tggagtaatt ttgaaaaaaa aattgattat tctattactt ttaattgttt gataccattt 240
 gtcattaaga tctccttcaa taggaacttt cttatttcca accattgaga gattaccctc 300
 gttggccaat gagaagaaag gcaaaatcaa agtntgtttt tggttttaat accccgtcta 360

gatatggaga cgagaaaaag gtaaaatgat taaaatcacg tcgaaataca attataatta 420
 caaatctgtt gtctagcaca atct 444

<210> 30720
 <211> 379
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30720

agctggtatg ataatggggt acccatcaca tgtggtacta ngtagggcgtc gggcgatggt 60
 gcacaacaag tttttcacat ccacaatgca cgcataaacc caccatcccc tgttgcccac 120
 ctccaactga gctcgcgtac tcccacgtag cccatattct cgtttctctc aacaccgggt 180
 ccccatcaat cctcccaagc ttccacaaca tccgagcaaa acaacattca gacagcacia 240
 gctatcacag ccaagcaaaa cagagcaaag gcagaatact ctgctcaaca catcaacca 300
 aatcacatgc tttctcactt aaagaccaca ggtacaattc ctctatcca attcgtaaac 360
 cgttggatcg actccaaat 379

<210> 30721
 <211> 158
 <212> DNA
 <213> Glycine max
 <400> 30721

actcacttac actggtgttc cttttttctg tgtgtttata atgatattaa ttagctgct 60
 attttttgag ggaacacacc actatttttt gtttgattca ctcaccaat atgggtaatt 120
 gatatatgga ttgttatttg gaactggaat ctttgat 158

<210> 30722
 <211> 373
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30722

agcttgatg tgcatgtggg cgtctatttc gactttccta tgctgtctct acatacataa 60
 aacagcccca ccacccaat ttgcaaaat catatatcat tggggcattt caccgagcac 120

ttgatgggcg catgttttga cataaattgc aagagaatgg gggcaatgtg gcatgcccc 180
 ttgcttcaga atacaacata ngcctaaggc cttctcattc aaataactcaa ctccacaaaa 240
 caagcatgga ttcagatgca aattgcttca cgaattntac aaaaaatgag caactatagc 300
 accaaaacac atcaatggag agccaaataa ccaagggaaa ttgcacttac ttgtggggag 360
 tgatttatag cgt. 373

<210> 30723
 <211> 1072
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30723

agctcggccg gaggcgggcg nacntcnagn actcgggacc cgtgctcgtn tgcgantagt 60
 caaataatag anaccacgt cgnctcnt acnacnctc cncnncccc cccaaggatg 120
 atgcagcagc gttttgcanc nnactagttn gngttgtaga aaactcnta ctctgcgang 180
 ancatacgtc antancnncg cntgtantcg acancagatc gacgtctacg atngacgata 240
 ncgacaacat cagcntatta cggagcgacc agaccgatct gtactngtgc gtcgcatttt 300
 gantcgacat ctgcantgaa ctaagccacg gtcctgngtg acatccatcc atgtancgta 360
 tatgtacgta ctaagntaca cataactaatg tccgntcgcg ttgataacta cganancgtg 420
 cgttgactg catcagactc atacgagcgt cgacntcatc tatctgtcct gtgnganana 480
 tcgtacactg ctcgatatgc tgctanagtc agtcgatagc tgcagtgatt acgcgtcgaa 540
 tgtactgtgn gaccngacga gtatgcatgc gngcatgacg cacacatact cctccgctcg 600
 ctctgntgcn tcantnaagc gtacgcgatg agatcagcta ngacgcantc atcacgcgaa 660
 tcatagtcgc gcatgcagat cgagcatagc tcgataagtc tcgacacggc tgcgacntat 720
 cgtgcactac atcgtctatg actgaagtcg gtgtaatcga tgactcatga tatcgcantn 780
 ancatataga tgatcggaca cacagntcta cgagtatgtg tatcgtgtca acatgcgtat 840
 gaacaagtgt caacatgcac nagacgtacg tctccgntgc gatgaatatg gatgactagc 900
 ctacgtctac gtcactact gtanagtcgt cagccgacac tgctatactc tnatagtgcg 960
 aagagtatcg catacacgaa cgagtatang cgctgcacgc acncgatccg antgngtcta 1020
 cngngctcct aacgtgatac gcataccgca gactctggcg cacacgtact cg 1072

<210> 30724
 <211> 391
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30724

agctntaata taattttttt caaattatta gngtatgtat ttatcgaata atatataacc 60
 tttatatgtt ggggggttaa agatgtttca gattccatgc tatcttcttt tcagttgcag 120
 ctttatgcc aatcaagtca ctacttgaat aaaaccaatg ggatgctgag cttcactagc 180
 gaattgatag gtaaagaata tagaatgtga tactaagtaa aaggatttca aacaaaagga 240
 taaagaggaa cggcacatag tgggcatttt cctaaataaa gtataaaagc atatgttctg 300
 aatgttntcc ctcataaaat attattgacc attgcatttc acaatntggt aatacctctg 360
 cttcctctcc attttctgat cactnttcta c 391

<210> 30725
 <211> 505
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30725

gatgccccac atagaacncc natgcgtgna gggtcantct atagaacctg caagctctga 60
 tgggtgtcgac aagacatcac atgtntgtca tcatcaaaaa tgtggagaat gtgaatgtct 120
 cccnncccc ttttcttcta ttcgtaacata taatactaca atgctgcctc acctgattat 180
 cactttgctt ccaatactat tttatactgc tccaccaac aatcctctgt actcacattc 240
 gctcaaatec atccttgaca ttcgcaaccc tctttctctc tgaccagtt tccgctttga 300
 tctcctacaa tctaattctc tactcactcg ctatgtcacc gtcgcgcat tccggcctct 360
 gcacctgcy caacctcct cgctcccgtc ttccgatctc gtccggaata aagcccgatc 420
 ccataacctc cctatactt atccagtccc aatctcgcta tccttgccgt accacctcac 480
 ctatcgtctc tcctgtcac cctcc 505

<210> 30726
 <211> 359

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30726

agctntgatt caaaattctg actcaccata aaccttgacc caaggtgaga atgccaatcc 60
 ttatcctcgg aagcaaaaaa agaggagaag aanatttcca atcaaaggaa aaaggagaag 120
 aaaatttcca atcaaagaac aagagaaaga aaattttccaa tcaaaggaaa aaaaggaagc 180
 aaagaaattc ccaatcaaag agtgggagaa agaaaaaaag aaaagaaagg aaattcccaa 240
 ccaaagaatg ggagaaagta aaaaagaaga aagctcctga tcgaaagaaa acagaagaaa 300
 tgtgcagaga ggtcttttga ccggacaata tctgaacaat acagaattgt caccaaag 359

<210> 30727
 <211> 466
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30727

gcttatgcgc atatttcctt acaaacgttc tcttgacaaa gacatttatt cgaanaaatg 60
 caccatata caatcaaggc agcttcgtca tctagattat ttacacgtac ctccaagggtg 120
 tatttgttac ttatatacaca cacatctcct tggctaaatt cacatacatg cataactcaa 180
 gcattttggg gtacaaaaaa ttgcacatgt gcacctcttg gcattttctaa tacctataca 240
 tacgcaaact ttatgatgaa tcttgactat ccacacaata aggtgctaca tttcatgcct 300
 ctttttcaag tttttgctac ctanagccgc atgcanaatc aagcatattt tcctttgctg 360
 actaaaattg tattcaaatt aaaaggtata ntttttgtaa tatgttntct tcacataaca 420
 tngcacatat ntatatatan tttttttttg tgagaacatt tgacta 466

<210> 30728
 <211> 399
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30728

agctgactta cactctgagc atanaagtgt gtnttctttt ntagaatgta tatangtgta 60

tggcaattag aatatattaa atgttcttgt atgttgacat gggtaatagg atactttcta 120
 cacatgcgcg tgtgcataaa tggattacat gagtttggtc taaatcagaa gggctagcac 180
 gacatttttg cgtaaataata agcattatct tgtaaaacta acttctanat gtttgttctc 240
 gcaggaaatg gccccgagga aacttgctc anagagatcc angaaggata aagcggccga 300
 aggaactagt tctgctcccg agtatgatag tcaccgcttt aggagcgctg aacaccagca 360
 gcgcttcagg gccatcaggg atggtcattt ctccgggag 399

<210> 30729
 <211> 408
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30729

ntacatctca ggctaagcgc atattcctga aatctttgtg ttgcatatag tgctaagtgc 60
 cacctactgc gctaagcccg gatgctcatt ggaatttgaa acttcaaatt gggcttagcg 120
 tgaggtagg ctaagtgcg gggctttaa ctcaaagtc atattggcat gctaagtgcg 180
 ccaaacaaaa atgctaaaat gaattagaac ttccataggt gggtaccttt acacaaaact 240
 tttgcttctt ttgctgagct ctcttctgt gtgtgagcat tatgctgttg tgctcaagt 300
 actttctaca tcttcttgca ttttaattccc atccaagtaa gtagtgcttc atttccattn 360
 tcatactgtg aaacttagga tagacgatgt cttgctttgt tagcttgc 408

<210> 30730
 <211> 313
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30730

agcttgaata tctntgatct accaaagaaa ctaatgagaa gaataaagat ttcttgtact 60
 catctgctac agacataatg taagagctan aagggccctc tcaagaagtg cagacacctt 120
 gagtttgctc aaaaagagtc tatgaattgc ttatctacat ggatcgatgc cntttatgga 180
 attgaacaga gtcattttat tcgatatgct aaattcttca ttggtgtaga atcttaaaca 240
 atgcttttct attttttttt gattgttaac aattcacgta atctcttttt gaacaaattg 300

caccctaatt tct

313

<210> 30731
<211> 489
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30731

ggcacacctt acggataana tacctattgt gntaaatcca catagttata tcaagtatct 60
aaatctgacc cattaagccc atgaacctag gtggctttgc gaataacaaa cttctcttta 120
ttaagatcca tatatttgta ttgagtttta gggctctgacc cgtgaactta gtaaacttta 180
tccatgaact cgtgaagtat ccatngaate cgcctaatat gtgtaagtat ttataatttg 240
gtatgttaaa gttatggcca atntacattg tgattgctaa tttgtagtgt ataaaatatt 300
aatatgattt agtgtgatag atcttagctt agaaaatgat ttcatttggt tcttcaaatt 360
ttatatatat tcaactttttt tttaaaaata acttataata aatactttgt tttcaatata 420
tcatgtgtca gggcgggtcca tgtatttgcg ggctttacga atcagatatg aattcttaga 480
aaaagtcta 489

<210> 30732
<211> 362
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30732

agcttngta ttaatcctaa ccttgcatgt gggctttttt ccacttacgg gagccgccga 60
tgggcccggt gctactgcct ctgagttctt tgctctcttg ttgcaccatc tcccacgcct 120
tgtggacctt ctgaagtgcc tccacgttgg tcttattgaa gctcgtgca atatcaggtg 180
tgagctttta ctctagtggg gctcctctca tagggtagcc aagctgtctt atagcaagaa 240
cgggattgta actgatgcaa ccccttgctc ccatcaaggg aacatatgga aatcttccgc 300
acgaaataaa agtcctgggt cttccttctt tcatcgaggg aaccagtcac agacactcct 360
tc 362

<210> 30733

<211> 174
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30733

attactacac tcgattaaca cgtcttacct atcnatgttc caatcataat gtttctcggt 60
 gcatccaacg ttcgtctgta acagtcaacg tttaaattcg ggcttggcga tctaacttat 120
 gggattggac ataaatggac aactaaatct gtgaattaca ttaacaggga ctta 174

<210> 30734
 <211> 364
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30734

tatctntgat ttcccgaaaa ctcacctcgt tttatctgtg cattccaaat tctcaaaaga 60
 gtcagtcttg tggcatatca nattgcatta cctccgtgtc tttctaacct ccacaatgtc 120
 tttcacatgt ctcactcca taaatatatc catgatccat ctcacatggg cgaattagat 180
 gaagttcaag tgaaggagaa cttgacatat gaaacatttg ctttgaggat cgaggatagg 240
 cagacaaagc acttaagaac gaaagagatt ttattgggtca aggcagtctg gggaggtgct 300
 ttacgatagg aggcaatttg ggaactagag attcaaattgc gagaagccta tctgtcttg 360
 tctg 364

<210> 30735
 <211> 302
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30735

gccctgaact gtgacctgan ctcaccctag aacgcgaagg ggntcaaaaa ttgttaaatt 60
 tcgacccgga ggatcgtaaa tcaacgcggc cctgtggcaa tatgaatctc tggggcgtag 120
 actgatttat acgttgagag gaccgatgaa tctgacttag cagtaacctt attgttgcc 180
 ctagtgtaaa atgcaagtgt tgatggactt atcatcgtgg ttgacgttct cttaaaaaat 240
 ctgataagcg agagagtta atgcttggct tgagctcagc gagttagtgt gcgtgggaaa 300

<210> 30736
 <211> 360
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30736

tgcttgagtg tgtagatgat gcaagtgact gagacaaaga ggatgagaaa gacaagaagt 60
 tctggtaatg ttgggtgcta tgggtgcaatg ggagtggcag ttgtggagag gtgtggcgac 120
 agagatctca cgtgacattt tgggaaccct agaggtaaga atagagaaaa acatttnata 180
 accaagaatt taaagcgcca gagaatataa agtgggagct acatattgaa caagagaag 240
 aacattctaa gacggttttt acaaaaccgt cttggaatga cagtcttcta aaacgatgtt 300
 cacaaaactg tctctgttga anaatccata tntacaaaga tgtcactgtc ttatatacta 360

<210> 30737
 <211> 187
 <212> DNA
 <213> Glycine max

<400> 30737

cctacttact tatactaacc caatcgcagc attaagcccc agttgttctg aaatgaagag 60
 gcactccccg tatatagtaa gtaccatccc ggtttcacct ttctagctgc cgttctctta 120
 cactttacag ctacgaaatc accttcaatc tactaaatta taccattttc tataaccatta 180
 gtcgtca 187

<210> 30738
 <211> 315
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30738

tgcttggtta ttatgtaata tgaatttcga catactatga aagaaggttc tgaattctga 60
 tgatacccat aatcttgta ggtgcccgct actgtcagaa tatatattcct tatgggtgcat 120
 taagttaatc tttcattatt tcaacttcaa tgtaaacagt gttattatca cgatgagaag 180

gtgcgattat ntcgatatcc tgatcgtttc tttagtttaa gatataatattt ttgttgatat 240
 acacttaatt tggttcanaa caagttattt ataataaac aattatataa aaaaactaac 300
 tgatagatta tcatt 315

<210> 30739
 <211> 368
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30739

tatagattat ctagttgccca ctcatgcaac catttctgtg tcaactgcac acaaagagaa 60
 ngtagtcttt attctttggt gtatccttct attatccttg ctaattgtta tgggaatctc 120
 tttctaccta ggcaactatg tgctgatcac aatatacata ttattgtaat gctcttaacg 180
 tattatcaat cccagcaatg gaacaaagct caagcgtgct gtaaaatggt ggcatcaag 240
 atatgggaca acttcaagac gttctgtgaa aggaatatct ctctttaac tgctggaatg 300
 tgacatgtga ttaagcaaag aactacagtc tgtgtctatg aggtaaaaat cagtcttgcc 360
 tcctttga 368

<210> 30740
 <211> 767
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30740

tctctaccct gcaactctact atccgcgcgt acancgcagt ancnntcta ctcantgtca 60
 nttttacctc cccacccccc cccnccaacg agagcgcgca gttggagAAC anctagntag 120
 agcactcctc tacgtactgt gcanactgan acatatgatc gtcaaacanc tgntgtaggg 180
 actatctnca gctaatacag tgatatctnc aatcatatgt gtatcatcac atctgtcgna 240
 ctatatacgc ctgtaaatgt ctgccgatca gcanggtact gagtattact tatctcagtc 300
 tagcaanact cgcgacgcgc tcagatacga cacaaccgc atcaatacag ttgagactac 360
 cgtcgtcagt cgacntaatc attatcgtgc tgacattcgt ctgcgacgta aaaactcact 420
 cacggatagg cgcacgcacg aagtgcaact caaggatcga acagaatgcc aaccagatat 480

ctcgatacac gacatgcatg tactcgccgc aaactaactg acgagacgaa gatcgtatac 540
cagatctgat gaagctgcga cggcgaaact ccattgacca gactcgcnac tacaggcggg 600
cgataatcgg tcgcgggacc attctgtagg gcgcaacaac aacgacgatg cgtntttctca 660
ctgagtgcga gaagacanac aaatcgtaac tacgtgaggg cgaactacgt ccgtcgacga 720
gtacatccgc atggagctta atcacctcgc ctagaactaa tccatcg 767

<210> 30741
<211> 302
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30741

ctgctttttg tcttgcgagt tgattntagc cttattttca ccttacttat tagtcaattc 60
aattaagaat gagaaatccc atagagaaaa atgtccgatt gattttccgc tctattttac 120
taaaagatga tttttttatt attatattat cttatacctc tttttgatta ccaatgtgat 180
tacttgacga ccgaacgggc gtaatttatt ttaaccgaag ttaacggata atacaattca 240
actttcggtg gatatttttt tattnttaag tcaagcgaga aatgacttaa gctaaatggc 300
tt 302

<210> 30742
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30742

gttgcatgg aattgcgaaa gcccactcc atcttttagga tntgtttctg ccatctcana 60
caaacaaatc agacgtaaca agacaattat agttgctgtt tgaatacctc actcactcaa 120
gtgtatcaca caattatggt ttttctctaa tgaaacactc ttgcctttta ccaactctaat 180
tccccttgag ttcttatgca attcaagaga ttatggccac aacagagAAC aattcaccaa 240
tatgtgtaag gtaaggctag agaaacancg aaaagggttaa ccaagaaaaa ggctaacaat 300
gttttttaggc acaaatgaac gaaacaaatt tcagacttta tgaattcaag taacaatcct 360
tcatgcaacc aatata 376

<210> 30743
 <211> 388
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30743

agcttgctta ttaaattgat ttggatccat atgacanaga acaaataana caagatttan 60
 aattganatt ggagaatgca agattgacaa tgggaatgag aatgtgaaaa ttatgtgcaa 120
 attgcttcct atgtgaccaa tttataggac ccaattntaa aaaagtttaa tgtaaaaaaa 180
 atataaaaaa ttaaaacata acatgcatcc aaaattcaca gagcaattgt caattgtatg 240
 caacgttcta aaattcatag agtaacgggc aattgtggca aattgtcttt cttttctgca 300
 ttctttctct ttntctttc tttctttctc ttcttcccc ttccaaaacc cacctcctat 360
 tgccctatct tctcttctct tctttctt 388

<210> 30744
 <211> 439
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30744

ntanagcaca acatcacaga atctaggtgt ccaacacccc tcaattaatg ggtnttctaa 60
 gtttgtgaag tgaaattgag aatgaggtaa atttgagca aactctcacc tcacacaagt 120
 ctataacatc aatctaaact tgctcaaact ggatntacac ctaaaattcc accgaatcaa 180
 aatttgactc ctcaacaccc aattttgccc tagaaatggc tcttggttca ctttggtcat 240
 ttgtttttcc ctctagcaca gcctaacctt tctcataagt cctaaatggc atttcaagct 300
 aagattaatt cactctaacc tctacatact accaattcca gaattggcct tccagccct 360
 caaaatcact ctntntcact cataacacca catnttactt tctaagccta ggttattcta 420
 cattcctctt acagtttcc 439

<210> 30745
 <211> 382
 <212> DNA
 <213> Glycine max.

<223> unsure at all n locations
<400> 30745

agcttggtat gtatatgtta caatgttctt aaatntctaa aaagttttta agaacaacct 60
gtctaggtaa atcttttcag aaagacttct aacacaataa gaaaagaaca gtttttcata 120
attaccttat acaccagcta atgatagaag ctctttcata ttagtttttt tcaaaagata 180
tttgtaaatt atgtataaac taacattaac ttatagaaca gtttatctaa ttttttctt 240
tttattctct ttttttagta gtacttctaa atacatttat ccaaataagac ccttaattatt 300
aatatatatc aacaatactt acatccaaat tattacttag tcaaggcttg aaattattta 360
tataaaataa ccagattaat ta 382

<210> 30746
<211> 645
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30746

aggatacgca ggtgatgcgg acacngctgc gtcgctgctg tacacatctc gggcgggang 60
antcgtgccc cactcagcat gagagatggt ggangccatg ggaacagccg ccatgtgcac 120
gatatcactc agcgggtatta cgcaacgcgc cgatacaaac atcgacacga cngtctgtga 180
cggtcgtgtg tctcagcgca aggcggttgg atatgtggac gtgtccatca cttatcaaga 240
gtgatctctc tgtcggtgca gaactatcag tgtcaggtaa taacacgagc agagtacata 300
cttggcgtga actagtactg gacggttaata cagcgggcca gacgattgtg cagtgtctat 360
ggccgcggac tcacatgtcc gcacgaaaac ggatgcgacg gttcgagtcg cgcgcatgct 420
cctgagcatc agatcagccc acagcatcac tggcatacat cgcgtggagg ctatcgcgcg 480
acgcttgaca atgtcgagcg ctcacacgcg aggtccgaac aagtacgact accgcgcgct 540
ctctcaacag cgctccacgg acgcaccgag cagaaatcgt tcgcgagaga tcggccggct 600
gaagttagtc ggcggactat ctctgggcca ttcacgaatg gaacg 645

<210> 30747
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30747

agctntgggtt ctcttcttct tcttcttttg gaacggtttc tttctgtgtg tcttcgtgcg 60
tttccgccat caccactcgg gaaccgtcgg ggcttggaag gggttaatgc gtcggtggcg 120
gaaatatgaa tgcggcgctcg ttttaaggtag ttcgagtttg gcgcacctgc agtgtgtgaa 180
tgtacatgaa ctgcttggct ttttgtttac gtcttcggag cagagaaaca actccaaaag 240
tcacgatat cgggcatagg tgggtgaacg tgtgaccggg cccaaattgt tgtgccgcaa 300
cacctgcgtt g 311

<210> 30748
<211> 377
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30748

taatgaaata naatagaaca gaagtcataa aaacaaacat ccatgatctc attaattctt 60
ctnccccatg aaatcctcac ataaatatca ttctgtacat ctcaattaca aagggttgtgt 120
cggagaataa taaaactaag atcaaaccga taaggcatca ctaattacat gtgttggaac 180
aatagagttc tcacacgcac tcaactgtcac tctatgtgag agaataacag aagatgatga 240
ccaaattgat tgagagaaaa tagaggggac ttaaatttg aaaaaaaact tctgcattct 300
catgcacact cttgcacact ctctgtttca ttgatgatca ttagtatttt tatccacact 360
gcatatgtag ctattat 377

<210> 30749
<211> 388
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30749

cggaaggatc anagcgggtc tgannagagg caaatttaat catcccactt ggacgaatga 60
gaaaactggg gcaaatggag agggtgagaa taaggagaa gcccatgtta tgactgccat 120
tcctgtacgg ccaagtttcc catcaaccga acaatgtcat tactcaacca ataacaaacc 180

ttctccttac ctactgcat tntatccaca aaggccatcc ctaaaatcaa ccacaaagcc 240
 tacctaccgc acttocaatg acaaacacca ccttttagcac aaaccaaaaa caccaaccaa 300
 gaagtgaatt ttgcagcgag aaagcctgta gaattcaccc caattccagt gtcctatgct 360
 gacttgctcc cacatctact tgataatt 388

<210> 30750
 <211> 442
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30750

aatactcagc ttgtcaggaa gctacctaag ctataaatag aagcatgtgt accacttttt 60
 gtaactttta tgaatgagaa acttgtgaga tacacttcaa agttccactt ctctcccttt 120
 cttccttcaa ttttccatgc cactttctcc ctctctcatt ctctctctct tagaggtgaa 180
 gcttctcctt ccattgctta ttctctagtg gatgacacat cctctctcct cttctccttt 240
 atcttccgct gaaactccat gcgtgaaaat cactattgaa ggaccttatt gaagctcaaa 300
 gatccagctt ccatagaagc ttctcaagag agcttncatg aagtggatc agatgacaag 360
 agtttcaagt aggtgctcct taaacctcca tttaatttca actttacctt ctctacatt 420
 ggtgggttctt cattatctcc at 442

<210> 30751
 <211> 360
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30751

agctnttcta tgaatcctct tacatgagaa gaanataatg aaatgagntt ttttttttaa 60
 actaaaatta attctccatt aactaaatat taattaattt atagatatca tatcatctat 120
 gagagaaatt atactaaaga ggctcttatc ctcttttggg ggggttattt gctctatcta 180
 gctcttggtc tctatatctt tgcaaatacc ttctcaatgg ctctgcatag tcgtcaaaac 240
 caagcgaccc caaggcccag catatgtcat ccccggtcac tgtcttactc ctttccctcc 300
 tgcacttctc cgacgcctcg ctgggttacia agcttatgaa ctccgacacg cactcttgca 360

<210> 30752
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30752

tgttgtcaag acaagacaat tgtgttcaga gttggaaaac tacatgttgt tggaagtatg 60
 aanatgaata attactttct ttacttcct ttattatttc tgtcacttga ttccataatt 120
 atgcatgtta attgataggg acttgggtat taaaggggtgc ccaagtccca catagagtag 180
 tatttaagtg cttgggttct ccccttaac aactagcttt taaaggtggg ttcaccaagt 240
 gcttgggtgc ttacattaat aatcctttca ccttttactc cctccattcc aaattgattg 300
 atgtttaggg attaaataat ccaatatatt actattcttt caagtatcaa atccaatgag 360
 atataaaaca tctatccttt atgcctctat aata 394

<210> 30753
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30753

agcttcctgt taaatacaaa acanaaaat tagacaaatt attatgaaaa aatngacgta 60
 aatgacatan attattagta acacttacca ctgcatgtct caactcgtca acatcagacc 120
 ttacagatgt taatgggtact gctgggtccc ggacctgagg gatatctgtc tctgggacct 180
 gagggaaaca ctctggatgc gtagcataac catctggcan aggatctgat ggctgggtccg 240
 atgtcatgaa tggatgcgaa atgcggaaga actagtccat gtagtcgttg gcacactgna 300
 cctgcacaac gcacatctca cctgctgcaa tcatatgggtc cgaatagtgc atcccacctg 360
 tgtgtatatc atcanacgac acccatgaat cgat 394

<210> 30754
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30754

cagcttgcac aactagagcc atcaaccaac tccagtgcac agaccatcgt taatttaatt 60
catcatgata aacattatac aaaagccatt cttgtggttg ctactagttt gtttcactac 120
atgcatgtat tttgtattgt ttttaacactt atgttagatt gtttcattat ttgtttattc 180
tgaagttgga tttatattgc cattcattga gaatatatat tttatttttaaaaataaatg 240
gtacaaaatg attgacaact gatacaaaat agaaatacat ttctttgtgc ttttgtgatc 300
aacaanaca tgtttccatg taaaggcatt tttgtaaaaa atacctanag cataacggta 360
tactcggcaa agagaggagg tgtttcaaca attntgtgtc ttgggttttc tttttt 416

<210> 30755
<211> 364
<212> DNA
<213> Glycine max

<400> 30755

agcttcattt aatccatgcc gacatctgtg gtcccatttt gcctccctca cacagcaaca 60
aaaggtacgt tetaagcttt attgatgatt attcacgtaa agcttggatc tactttttgc 120
atgaaaaatc tgaacaaaat actgtgtaca aaagcttcaa agcctgtgtt gaaaaggaag 180
ctggtatcta aattgtttgt ctaagataag atagagggtgg tgaattcacc tctaaagagt 240
gtacagaatt atgcactaat caatgtatct ctaggcaatt gacgggtgcc tacacccac 300
aacagaaagg agtcgccgaa cgcacacacc gaactatcat gaatgttgta cgagctgtat 360
taca 364

<210> 30756
<211> 475
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30756

tgaacgtgat gaccntgca aactgaacca taacgccgcc gggtgaaatc aaacctgtag 60
agatttgaca aagctgtatg caacgggaac aattttgttc ctaatgaaat ctctctaacc 120
aagcccaaga tagaggccta tctaagattc tactcatgaa tcacagcgct gatgctgaca 180
agattatttg gcatgagggt gtctatctct aaactgggtg ggatcagtgc actatactct 240

cacttcaa at aatgactgat tgtccatcga ctgcggagac cttattgaac cacctgagat 300
 tgcgacatgc tggattgacc cggaagagtc aatgcggtcc cagattatcc tfgctatgta 360
 ttcactactc tatgagatgc gcccttgctc aaaaattggt aaaccctatt agagggaaca 420
 accgagaacc tctctgtgaa cgagaatata tttcttaagg gccggaagg tttct 475

<210> 30757
 <211> 395
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30757

tgctntatat gattatgatc atgtaacaaa tcaaagaaac caaccttgat gcatgcattt 60
 ttgttctgaa acgaaactaa agaaacaaag gaaagggaga aaatagaaag ctaagttcta 120
 agatacaaaa tgcccaaggc atttgtcggg gaattcgagg ggagtaaaca ccagacaaat 180
 ttacaccaat gagccatgag caaccacata aggaattta acaccacact ttaacccaaa 240
 accttaaggc tcaagtttat gggctcttctc cttacttata tgggtgctcaa cttttcaact 300
 tccatcctat gtgtgctcaa cttttatggg agcaaaagaa gaagctccat gctttgtcat 360
 ccagtcaaca cagtcaatgg ggattcatct tcata 395

<210> 30758
 <211> 325
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30758

tgagctctta gaagacatgc ttcttcatta gtgctnattc ctctttcttc acaattcagt 60
 cagtttaaga tctcttatag ctgtcagaag gtctcttaat gagctcattt catactgtgt 120
 gcaagaagag gatagactga agcacgaaag gactaaaagt gctcatgtag taagtacttc 180
 taataaccag ggccaaagag aaaggactga cgagcccaag aatgaaacta ccaatgggtcc 240
 aacacaaaag aaacaaaatc aatgtgacaa ctgggtcttt ttagtagcg ctgacattgt 300
 aagaagaaat gtccaaatat cattc 325

<210> 30759

<211> 390
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30759

tatctntcat cttattcctt cacaaataac ttccataatg cataaaccta gtagaactac 60
 ccatcatatc tcccagaacc caataccac aataatttat gtgagaagaa gtctacccaa 120
 acctgaaatt tgaagtccca caacgtagag gtgcgcttca cgactccgaa aatggcttcc 180
 ttttgcgatt tggagcagat atggtgagta aagtttgag ctttgatgga ggcttcagga 240
 gaggaagaaa gggagaaaaa gcaacgtgag ggagagggaa tagcttctga acttttggt 300
 gagtgaagag agatgaacgt ggcttttagt ataataaggc ttccttntt tattttttta 360
 caagggtatg ccacatgtct ccttttgagt 390

<210> 30760
 <211> 495
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30760

tgacaccgtc ttgagccgtt ctacnacgtt gacaacattg tgatcagccg tcaanaacat 60
 atagtcacgc tctggctata caacttttaa tgtgaatatt aagggatagg gctattttct 120
 aatcctgctg cggagattat gactctcgcg tacatatgag aagtcaaact cacgggtttt 180
 ttatatgtgt ctggcgacag actcaatgca tatgtccaac agagctaagg tctccattg 240
 gatggggaac aatcaaggct aacacgagct ctgtttatgc gtgttttact cggtgcgaa 300
 actactgacc ggggacttga ttcttgcag gaccactgtt aactgacca ggcttagtag 360
 ctgctctgag ggcagatttg aaggcggcta agtttaagta ctaatgaagc gcttatgtaa 420
 ccccgcgca taaaaatttc tctggcgctc ggggtgggta taccgattaa cgggtttggc 480
 accatggaca tgcac 495

<210> 30761
 <211> 335
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30761

tgcttgcat aagtctatac cgctgattta atgtaattag agcctataga tttcctttct 60
cttttgtttt ctgaaatcta cctcattaaa taaacaaaga gatcttggtt catctgttct 120
tgcagttcca ccttttctca tatcattttg catgtttttg tttctttggt cttgcttggt 180
atagatatga gggtegattc tttgaggatc ctaacaacga gggtttgaca atcgattntg 240
atagagatat aagccaaacg ataaacgagg aagaggaaga ggacgtcctg tcaccagagt 300
tggagagggtt ggtcgctcac gatgaaacgt gaatg 335

<210> 30762
<211> 325
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30762

gcttctccct atttgctatc ataggggaag atgtgaagaa gattttgggt cagcccctta 60
tgcatttctc tctctgtcga atntgctgag gaaaattatc ttcgtgaaga aaattcaagc 120
cgaggcgctt tcgtaacgtt tccgtgagta attacgcgaa gattctcgac cgttcttcaa 180
ggtecatcgc tcgttcttcg ttttcttcag tcttcaacgg gtaagtacct ccaaccagct 240
tttcatttca ttctatgtac ccgtgggtgt gcacattctg tttcatgtat tagtattccc 300
gttctcattt gctttatata ccccc 325

<210> 30763
<211> 389
<212> DNA
<213> Glycine max

<400> 30763

tttcttcact tgaaattaag ttattcaatt atatgagttc ttgatttaat cccaatattc 60
tctccccctt tggcatcaac ataaagccaa agtgtgtata gagacataaa atcatacaca 120
aactcataat catccaagca ttttaattcca tacaacaagc aaggaggaca ataattcata 180
cataaactaa gcaaggaaga taataattca tccattaact ataataaagc gtcaaataat 240
tagaaagtca tccaagataa ccgaaataaa aagactaatt tagagagtaa tataactaata 300

agtgtatcaa atatgtcata agacatcaac acatataaca aatcacttgt ctaagtcact 360
agcatctaga agttctaatt ctcttctaa 389

<210> 30764
<211> 327
<212> DNA
<213> Glycine max

<400> 30764

cgcatgtcat caagtaataa tccccggacg aatttatggt atgatatttg cccctcttta 60
cttgccctctc atcggagata agatgaaagc aaacatagga cactgatctc gtccgtcctg 120
ccgttccccgc gatgacgact cacggctcta ttccttcggt tttcttctgc atacaacaaa 180
atacgaacta caacgagaac aacgactatt atgtacatat acacatatac acatatccgg 240
cgaaggaacc gaaccagaaa acaccagaat tacgggtttc ccagtcacca gaagcttcgc 300
gcttgacaat ggaggacaca tgaatag 327

<210> 30765
<211> 377
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30765

tgcttcatga tgatgaatca agattgattc aaggagtnt gatgataaca aagatgatga 60
caaaaagccc aagagaatga gttcaagatt gagtcaagaa cacttcaaga atcatgagaa 120
atttgatttc aagattcaag aatcaagttt caagaatcaa gaatcaagaa taatcaagtt 180
gaagattcaa gaatcaagaa aagactcaat caagataagt actaaatfff tttttcataa 240
cattgagtag cacatgaagt nttcacataa gcttttacca aagagttttt actgtctggt 300
aatcgattac cagtntactg taatcgatta ccagtagcan aagttgtnt caaaagcttt 360
cagattgaat ttacaac 377

<210> 30766
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30766

tggggggacan aaccaaacc caagagagaa attaacaaaa cctatatgag aggggagcaa 60
ttatattcct aaagaattct gtcctaacaa aaccaaagac agagtcttag caagtgtagt 120
tctcatgagt caaagtgccca aggttagcaa gtttatttac acatgagttt tcctctctat 180
aaacatgtga gtaacaaaaa actatatattt tacaaaaaan aataaacatt ttcccattta 240
ctatggagac attaaggaac caacaaagga ttgctgaaag cctgaatgac caatgaagag 300
tcacattcaa tccacaaatt attccagcct ttgcacttca ctacttctaa ggtgatgcct 360
acaccttagc aaaaaaattt gtttaacccc tata 394

<210> 30767

<211> 346

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30767

gagggagaga tagagagagg gggggacatg atattgaatg aagaaaaagg gagagaattt 60
gaactttgag ttgtgtctca caagactctc attcatcana gttacaacta gtgttacaca 120
tgcttctatt tatagactan gtagcttctc tgacaagctn tcttgagaaa acttccttga 180
gaagcttctt tgagaaaact tccttgagaa gctagagctt agctacacac acccctctta 240
taactaagct cacctccttg agaagcttcc ttaagaagat tcctaaacaa gttagagctt 300
agctacacat acctctctaa tagctaagct cacctncttg agatga 346

<210> 30768

<211> 467

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30768

cgctctanat ctacatttga tggtttgtat ttatgaggag gagggatatat tgtcatttct 60
tgctataaga gtagtgtccc actggtaaaa ttaactttcc aaatggttgc cttcgcatga 120
atggccacga ggaagcttgc ctcaaagagg tccacgaaag acaaggcggc cgaatgaact 180
cattccgctc cggagtacga cagtcaccgc tttaggagcg ctgtacacca gcaactgctt 240

caagccatca aaggatgggt cgttctccag gagcgacgcg tccagctcaa ggacgacgaa 300
 tatactgatt ttcaggagga aataaggcgc ccgcgggtggg catcactggg tactcctatc 360
 gccaaagtta tacagatata gtcctttgag tttatgcaa tgccttgccc acagaaggcg 420
 tgcgtgacat gatatcctgc gttacgggtc agtggatccc gttcaag 467

<210> 30769
 <211> 384
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30769

agcctttatc ttattctcta acaccatgct ttccatcctt gattcgctccc tctcggcctt 60
 tctaagcttg atctcgttac tgctgcccc aagagccctt cggaacttgt tctgctcca 120
 ttcttcattt cgggcccatt ttgtttctcg ctctaacgct tcaactgtgg tcatgttgat 180
 atccttcaat tcatcacact cttttttgac cctagtgact ttcgtcttca gttctctttt 240
 caccactctt gtctttttga gttgtacttt caaagcttgc acttcttcac tttccttagg 300
 aatttcagcc tttntccac ttagacattn tagctntggg agccaagtca tcccttgctg 360
 tctagacttc aaccacttgt gata 384

<210> 30770
 <211> 557
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30770

ttgccacaca tagcattgtg tgtgcaatta tatatttgc cgtctccag agagacgtga 60
 cacgtgggtc ataccttaga taacctanag cacttgatt cactatagat tatggcgtag 120
 atgcgtatct tgaaccatat atttgacta ctccgcgga tctattgaac tattgaatag 180
 aattcagtgc tcggctgctg aatcactcga taattcggcg taaagaacgc cgcgtgctcg 240
 ttatatattc agatgtaaga ctgaactggc cagagtgttg aagacttctc ggtcgtcctc 300
 gcgcacgcac atgcgggac tcataaatgc tgggttgaga tctctgcac ttattaaaat 360
 atgtctacgt cgagcatgcc taatatctcg tagccaatag caggtgaaa agacatgcgg 420

cgctcagagg acgcgcacat agtgagcgtc tattctggta gtatattata cgtgctgcat 480
 atcggcacat aaatgtaata gagaccagtg gcgcagtcgg accggcacia gtactcggca 540
 tgatttggcg tcaaacg 557

<210> 30771
 <211> 320
 <212> DNA
 <213> Glycine max

<400> 30771
 agcttcaa at ggaagagaga gaccgatcac gagcacatag catggtctta aaagaagagt 60
 tagccgcttg ctcaagggtcc aaaaggaact tgactcaacg tttatgagag atagagacca 120
 gcatgttagc tatcatcacc aagtaccaag aagaactaag tctagccacg gccacaaagc 180
 atatggtggc ggacgagtat gccaagtct acgcggaaaa agaggctaga ggaagggtaga 240
 tcgactcggt acaccaagag gaaaccatgt ggatggaccg atttgctctt accttgaacg 300
 ggagtcaaga acttccccga 320

<210> 30772
 <211> 350
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30772
 tgtagaatgg ctagacatga tacatgtcag gggttggttt ggttcaagga taaaaggat 60
 gcccacatt atttccatga cacanatgca aaaatgatga tttggaaatt ntatgcaaaa 120
 ctggtcatgc atgcacctat gcggacactc aagtgtcaaa tttttatggt catgtgatgc 180
 tagggctcaa gattcatttc ctctatttta gatcaacca atgtttccaa aatatgttct 240
 tttatcaatt tgtgcattca tccgagtcca ttttgggtac tcgggagaat nttcacagca 300
 ttcacccttc aggtgtgcac acattttttt ttcaacaact agctatgatc 350

<210> 30773
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 30773

agcttgactt gatgtaagac acatcttctt caacctttgt cattcttgac tccatntcat 60
tgaagcactt atgcacttgc aattccaaag tatcaaact ctcaccaaca aaggtttgaa 120
gaccatcaaa cctgtccata atctttgaaa gaagagatga atcttctcct tcatgtcctt 180
cttcaccaac atttctagca ccttcttca cccaagagcc atcatgctcc ttacataac 240
caaaggatgc tatgactgaa gtgcctataa ggaatgatct cttgattgga acacaagggt 300
cagaatcaag agggatattg aagtgttgaa ggaaaagggt aacaagatga ggataaggca 360
atgggtcatt caatcgcaat gccttatgca tgcgatattc aac 403

<210> 30774

<211> 402

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30774

tcgatggaga agaataaagc gcgagcaaaa tagggctcgc aatctaatat tntacaatgt 60
atgtacaaca tcggttatca atacaaaacc gatgttaact aaatgatgtt aacattaaca 120
tcggttttct acaacaaacc gatgttaacc tatcttatgt taacatcggt tnttctaana 180
atcgatgtta acatactgac tttaacatcg gttattcaaa aaccgatgtt accagtttca 240
tgttaacatc ggttttttaa caactgatgt taacataagc taattaacat cggttttcta 300
aaaaaccgat gttaacaaat tcacattaat tacaattatg ccaccatgtt aacgttaaca 360
tcggnnttga ggaaaaccga tggttaaacgt acgatgttaa at 402

<210> 30775

<211> 314

<212> DNA

<213> Glycine max

<400> 30775

acatgtcgct gggttcagcc ttgccgtctt gagaagattg taacctgctt ttgctggcat 60
gttatactca aggtttgcat atttgagtgt cgttggagat ctttcttacc ttgcttattt 120
tatcaggggc ttcgtccaga aagatgggtgc taccatctgc attcatggca cagaggataa 180
agtactgggg ttgaaagcca aatatgagga cgttgcatgg ggattttaga gactgagat 240

<400> 30778

tctccgaggg actcttgatc ctgttacttc catattatct cttcatatct tctcttacac 60
tctcaaagta tacaaatatt tcatttaaac ctcataaatc atggattctt gttctatttt 120
ttaatgctgt gggtgcttgg aatcatgaaa tatcattgga ctttagttct atgttgcaaa 180
agaatgaaac tacatcaatt tgaattttga tctaagacct tgctcagttt ttatttaaatt 240
cttgagtagt aattgttagt aatcttagtc aatttttacg ttttctgtgt ctctagatcc 300
cgtggtaatg tttttggaat tgctgagcac tctccaatg ttaatcattt tcgtactctt 360
ttctaaacac tatgtgattt ttatatgtat cactacttcc tatattatac agtttttata 420
ttgatacacc atgca 435

<210> 30779

<211> 412

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30779

agcttcctct gtgccatttc ctgcgaaggc aaacatttgg aaagttagtt ttaccaagaa 60
atgctactct tanaacanaa atggcataca acctcctcca ataaacacaa acatcaatgt 120
aaatntagag caaactcatg cacatacttc cttatgaaca ttcactcgca caagatatcc 180
ttctatctaa gaaaaatgca cccatgcaca atcaaggcac cttcattacc tagattatnt 240
atatgtactt ncaaggtgta tntgctacct acatcacatg cacttncttg gctaaatnta 300
catacatgta tactcaaagc attttggcta ccanaaattg cacacgtgca cattctggta 360
tttccaatac ctatgcatat acaaactntg tgatgaatct tggctatcta ca 412

<210> 30780

<211> 321

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30780

tataagcagc acanatggag aagttgttca attcatggca tcttctatc taaactcaag 60
ctttgccgtc aattaacct aagcacaact tgatttatgg ctttgaactt tgaaaattga 120

ataaaatcca ataaacttat atttcgattc taacaaacta acatttaca ttttaaaaaa 180
taggttctaa ccgcacaaat ggaaaagttg ttcaattcat ggcattctct tatctaaact 240
caagtttttg ccgtcattaa cctgaggcac aacttgatta ggactatgaa ctgtgaaaat 300
gaatccaccc aattacttat a 321

<210> 30781
<211> 401
<212> DNA
<213> Glycine max

<400> 30781
tttcttctct atgaagcttc ctagtctata aatagaagca tgtgtaacac ttgttgtaac 60
tttcatgaat gagagtcttg tgagacacac ttcatagttc cacttctctc cctcttttat 120
tccttcaatt tcgtgctccc ccctctctct ttctctccct ctttcttttc ctccattgaa 180
gcaccccttc aagcttctta tccaaggctc atcttggtgg tgaagctcct tcttccatgg 240
cttattccct agtggatggc acctccgctc acctcttctc ctttgacttc cgtgcacatc 300
ccatggtaga aaatcaccat taaaggacct cattgaagct catagatcca gcctccatag 360
aagccccaca agtaagcttc catcataagt gcaactgacct t 401

<210> 30782
<211> 203
<212> DNA
<213> Glycine max

<400> 30782
tgactatgcg agttgattta gccttagttt ctctatattt attattcaat tcgactaaga 60
atgagaaatc tcaaagagaa aacgtccgat tgattattcg ctttatttcg ctaacagatg 120
gttggttgatt atgatattaa ttgtttacct ctattttgat tgccaacgtg gttacggcac 180
gaccgatcgg tctgattttt tta 203

<210> 30783
<211> 374
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30783

ttgcttgttg acacgcggag atntacgtca tcttttgtgc tcacaagatt tgtcatactg 60
 acatttgagt cacgttgacg ggcggagata ccctagtggg tatccgtata aacattcttt 120
 nttgctgtct gtaaaacgaa aagcctgata gcatgcagag actaacgtcg tcttctgcgc 180
 ccacgtcaa tcgcggccga caagcccgtt gacacgcaga gatttacgtc attttccgcg 240
 ctcacaagat ctgtcatact gacattngag tcatgctgac ggacggaaat acccaagtgg 300
 atatccgtat aaacattctt tnttgctgtc tgtaagacga aatgcctgat agcacgcaga 360
 gactaacatc gtct 374

<210> 30784
 <211> 233
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30784

caccgcttta ggagcgttgt acaccagcag cgcttcgaag ccatcaaggg atggtcgttt 60
 ctccgggagc gacgcgtcca gctcanggac gacgagtata ctgatttcca cgaagaaata 120
 tggcgccggc ggtgggcacc actggttact tccatggcca agtttgatcc agaaatagtc 180
 cttgagtttt atgccaatgc ttggccaaca gaggagggcg tgcgtgacat gag 233

<210> 30785
 <211> 591
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30785

acacccgacg cggcatacag acacgcgaac tggagaacag aagcacggac aganaggact 60
 cgcgaaaggc gaaaaacaca aaacggagng gnattggaag cccgtggaaa caccanggcg 120
 ananaggcna aancacacgg gaacccgcgg aaacgcacaca gagcgcacgg aggcaagcgt 180
 tgctgancga cgaaggaaca ccgccacaaa gggacgaacn gcagcaacaa gacaagaccc 240
 accgcgaaan ccacaacaga acgacacacg gggccaaacc aggacacgga cgaacgaaga 300
 anganagcgc cggagaacgg agacggcgga agcgacanaa cagacgcaca cagacaccgg 360
 gcgagacacc acgaaaccga agcgagagcc ggccgacggc aagcgaaacg agagacaggc 420

caagttggtc agtgggaatat gatgaacaac gtcagctaag aatgaacaaa taatgttttag 240
 atgtaattat tttattgaaa gataagaaaa gggaaaatta cttgtttcca acttactcta 300
 attctgggaa atctaattgca gaaagtataa atggaagtac tgtaagatat tgggcatgta 360
 tcaacttctc 370

<210> 30789
 <211> 405
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30789

ttgcttgctg agtacaactc ataatacaat tattaatgt tttttgagaa aaaccaaatt 60
 catttagcgt caatttcaag aagccccaat caactttatc atagactttc tgcaaagtca 120
 attnttagcc taagattccc ttttttatta tgcattgtgat gggcaatctc ctgagctatt 180
 atagcattat cagatggatc tctattagga atataaatgc tttgcaaagg gccaataaga 240
 ctatcaaaat gaggttgaat gcgattaaca agcacttcag agataatttt gagatcgaca 300
 ttgcataact gatgggccta aactctttta aggaagaagg gcaatccact ttggggatag 360
 ttacaataag agtttcaacc aaacttggat tgatggagcc taaag 405

<210> 30790
 <211> 453
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30790

ntcatgggtg cgagaggcca tgtgaagcta atattcanat atacatgcaa gacctctcta 60
 ggtgcgagtt gtggcaaatt ttaatttatg ggttaaagtt agcttgagaa agggattatc 120
 aaaacatcaa gatataatca tattccctta cggcaattaa tctcttgttt atcaagagag 180
 tttagctcaa ttagttgaat aaaatgatgt gttgttgtga atccttagta cctatcgttt 240
 gattgctaca aattaaataa acaaattctc cgtctaaagg tttctgttgt ggtgatccat 300
 gtgcgagttt ggtcaaagct atttatgaca gccatcggag taaaggccac atttactggg 360
 ttcatgtgct gatggaatcc catcaggtgg tagatcgatc ttatggcaaa tcatgggtta 420

aactttagag tcttcataac atggtgccta tat

453

<210> 30791
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30791

atctgatttt ggattggcaa agctaaatga tgaggacana acccatntga gcaccagaat 60
agctggcact tagtgagtgt ccatacattt cctttnttaa ccatttgcac tgatcattac 120
tgagctatat gttggatcca ttatcatact atattataat ttcgccaagc ttgatgatgg 180
ggataaaacc catttgagca ttaaaatagc tgacacttag tgagttccca tacatttcct 240
ttttaccatt ntgcattgat cattattgag gtatatgtta gacacgtaca ttatcataat 300
ataataataa tcaagaaaaa caatagacac catgtattga aatacctcat gatcaaatnt 360
anatggaact tacact 376

<210> 30792
<211> 283
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30792

ataatatagg atgattatgg atccaacata taccttagtc atgatcaatg cagcatggta 60
acaaaggaaa tgtatggaaa ctactangt gccagctatt cttgtgctca catgggtttc 120
tgcctcatca ttaagcttag cctagtccaa atcaaatac ttgggggttga gatctttata 180
ccacaacaca ttattggcct tgatgtccct atgaacaatc ttcattgttg actcttcacg 240
aaagtagcca aacctatagc gataccaaca cgaaatctat gct 283

<210> 30793
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30793

agcttaatct tatggcttgc ctccagactt cactccccgt accgctccag acgatttgaa 60

ccaagcccct accttcgagg ggcaactccc tccttatgcc gattatcccc tgcaagaaga 120
 cgatgaagaa gatgcccgtc taggcccctct actgcccctc aaggatccgg ccccccata 180
 attgcccctaa ccaaacaatag tccgccatgt cccatctcca cccgcacccg ttaaagaatc 240
 tgttccccttt gcaaaagata gggaaagatt gatttacttg aagagagggt gagggcggt 300
 gaaggcctcg acaactaccc gttctcggt ntggcggtatc tgtgtctggt acctgacatc 360
 gtcacccctc ccaag 375

<210> 30794
 <211> 810
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30794

gacccaacgt acctaantnc aagcacagan acgtacacag acagngntat cgacttaaa 60
 ctaatcnant cctactanc nanccncaga gagcgngat tggacgcat taggtagana 120
 ngccattgat aganacgtac aaccnaccga cgatcgacag cantaactat aaaacgatgc 180
 gatcgagacg tcaactacgta tatgtgtcac acaacgcata tcaactacca taagaacgca 240
 ganagcgacg cactatatga gacagcgagc ttatgaatat catcagcnga gcagtacgca 300
 ctcaactnct gacgcanctc nnatgtagac ggcatgatat acatcgacgc acatgaatga 360
 gatacagaca cgcatgaacg cacacgagaa gcggagnang atgatacgaa cacagccgac 420
 tacaactgag aggcctgtag gcacatacga cagcatacac tgcagcaggt agtctgcaga 480
 aacgagacag agagaagata cacgaagata gtcaacttaa cacagcatag gacaatacca 540
 acagatcgag catgaaaata acggctggat gcacaacgga gctgacgaca gcgccaacag 600
 acgactgtga tgcacgagac gtacaaacat caggacgtca ctgcgatca cacctcacgc 660
 agagtaacaa ccctgccaca cgagtgcga gaacgaacac gagcagacga ccgacgagca 720
 ttgtgtctta gggagatcgc acaacgagga cacactgata cgacatagac ggagtacaca 780
 aactgatctc actagacaca gccaacggcc 810

<210> 30795
 <211> 326
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30795

agctttttatt taataaaatt aattaaaaaa ataaggattt tttttacgta aaggetatag 60
atacaaagct tcacactaaa aaagaaacat ctcaactacg ttgacaaccc tcctcctagt 120
gatcacaagc aaaggcgtac aattttattca aataaaaaga agaaatagat gaccaacact 180
acgaaaagaa gtcttgtatg atgtctatgt taagatgggt atcgaaaagc ttcctcgggt 240
taagtagtgg tggcattttc gtaaacaatt ataacttttg aaagacgggc attgcanaac 300
cgtcttttaa acaacttttc aaagat 326

<210> 30796

<211> 480

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30796

gcgcttgac cctgttgan ccaatctaga cgtgacacct tggcataagc ggcctcagga 60
tgtccagatg gagaggggtt tcagactggt gcccgcatac gtttgacggt ttagatctgg 120
tcacaagaag tgtggactaa cctacttcca cgatactttg gattaatcag aaactatatc 180
acgagctact ggcaacacac ataattgggg ggcagtgagg tgcgacacag caccatatct 240
gattactggt gcattggaca cagcatgac cccacatatg agtctccgac gaaagcttac 300
acccaacga ctgacctctg cctggatgca caatctatct gcattgacaa tgacaaagga 360
tggtttctgg ctaatatcgg catttaacac gcgctcaatg tagtcactcg attaaaatgg 420
gtctcttttc cggaatctaa agggacctca tagctagata acacttccga ataaagatcg 480

<210> 30797

<211> 413

<212> DNA

<213> Glycine max

<400> 30797

actatatgcc tcgggcacac aagctatctc catactggct cgatgagatc aatataatct 60
cgcaaccttg tcagcaaaga actcatgagg ttataatgct tcactctaac tgactcacca 120

tacaagacca tttgctcttg ctgtgcatgc aatctggagc aattgaacac cctgaagcct 180
 atgctgcaga catcaacaaa agacctgctc tacctcaaca acaaaatctg ccacaacaga 240
 aaaataatga cctctccagc aatatgaaca tatccaggag gaggaatcca ttcaacctta 300
 aatggcggag ccgtcacaca accacaacaa caagcacaa cctatttcta aatgctactg 360
 gacaagaagc catatgtcca tcaccatcca cagcacacaa cacagcacag ccc 413

<210> 30798
 <211> 238
 <212> DNA
 <213> Glycine max

<400> 30798

ctctgcaggg aatctaagtg tgaagcatgc tattctgcac acgattgtag ctgccaactg 60
 ggtaccact aatcatactt ccaactgttg cacaagttcg agtaaatttc tgtatgctgt 120
 cggaaccaca tccaaatatt atctggaaac tatatctttg atcaaactgt caaacattca 180
 taatcttttag ctatcaaata agccattgcc ttccctactg tattgtgtgc attatgac 238

<210> 30799
 <211> 401
 <212> DNA
 <213> Glycine max

<400> 30799

caagctgggt tactatttct gcactaaatt gttgtgtgg taatcaaag cagatgcaat 60
 gcagtagaag ggggtaaaga caatatatta ctacaattat atgaaattga gtaggtaata 120
 ctaagaatag aatattagta gcatgaccga gaataaaata gccgttgtgt catataacat 180
 aacaattgtc tcacatacag ggaaaaaaa tactccaacg ccatcattag ccggttgact 240
 tattgtgtgc ttttaataaaa tgttgcccat ttcttttaaa tgtggtgatg atgccgatgc 300
 caacagtgtg tattatcaac cactgcaagt ttcataattg taagttagac tgcaaaaata 360
 aaattcagtc caattgttct gaagttagtc ataccatgga c 401

<210> 30800
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30800

actcaagctg atgcattctt ttctgcaga aggcaacaca atattcaaga cttataaaaa 60
tgatatgctg gtttggagaa gatgcaatct atgcaccata ttaaaccac aactcaacct 120
tccaccaac acatgttgac cctcctcccc aaattgaaat caaattcttc gtataatttc 180
tctaactctt ttctgcgca gctaacaaaa acctatatct aanatatattt aagaatgtct 240
gtcactacag gacatacatt agttacttag attacacaaa caattgaata atgagcgctc 300
taccttcata gtggtttata ttatcctttc tttattttac aacttatatg aagatggctt 360
tcttgtttct ccataaccac ttcattaagc atttgaacc ataactcttc 410

<210> 30801
<211> 316
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30801

agctngatgt agatgtatca tatatgngnn cnnaacaatn ggttnttgag tgcgtggtta 60
tgttcgtgag tggaattatt tgcattccat cttgcaatgg aagaatgagc agattttatg 120
tatgtgtttg actgttttgg taaaatgtga tagtctctta tggacaccat tcacctattt 180
gtgctgtaca atatatgagg atctgatgca ccaatttcat atgcaactgc caagtcactt 240
gaactgttgg ctgcccagag gagcttcata gcttgttcca aagctcctac ttctcaacct 300
cctccatgag tctgat 316

<210> 30802
<211> 364
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30802

ctgagaatgg ctagacatga tacatgtcag ggcttggttt ggctctatga taacatgtat 60
gcctccatt atttccatga cacaatgca taaatgatga tctggaaact ttacgcacaa 120
ctggctcatgc atagcagcct atgcgcgaca ctcacagtgt gaatataatt atggctcatgt 180
gatgctcggg ctcaagattc gtttcctcta ttttaatcgc cccaatgttt ccaagacatg 240

ttcttttatac actttgcgca ttcacccgag tccatanccg gcgtccggtg aaacatcaca 300
gcattcaccc ttcaagtga tacacgtttt ccataaattg tttatgatca atgaatttgt 360
ttct 364

<210> 30803
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30803

agctntgttt ataagctcga aaagacaaga gtggtgaaag agaagctaaa gacggaagtc 60
actaggtgca ggaagaagta tgatgagctg aaagatatca acctgacct ggttgaagcg 120
tcagagtggg aaataaaatg ggccctgaaag gaagaatgga gcaggaacaa gttctaaggg 180
gctttgtggg ggcagcagta atgtgaataa gcttagaagg gatgaatcaa ggatggaaa 240
catggtgtta gaggataagt taaaggcttg tcagaggtcg aagagaagtt tgacagaaca 300
gctgagcaaa atagaagaga atatgttgat aatcattgat caatataacg agaaggtgaa 360
cctagctgct agtcatggac atatgctgga aggtatc 397

<210> 30804
<211> 478
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30804

ttatcatcat tggatttggg gtatcaataa aatgggtatt gttgtaatgt tattttgtac 60
aattccacct ttgtacaatt aaaacatcgt ataaatattt tgtaatgtaa ttattgtgta 120
taattttaaa ttgtgtaaga atttactgcc aacgatttgt cgtatgattt gtgctactaa 180
gatccattgg anaaaaataa aaatagaatg tggaagaagt tttgaatctc ccaaagtaca 240
tctgacatgg atacaaactt gagtagacgt tgctttgcta catattgtac tctttcatta 300
attagaagtt tcatactagc ttccactctc ttcactcttc tccaatgaat gtaagaagaa 360
taaattatat aatggaacat ttgacattgt tgcttccttc ggtatcggcg atatatatag 420
atcgattctg aaaacataag tctgtgtatt atcattagca tatatgatcc ttatgttc 478

<210> 30805
 <211> 337
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30805

tttcttgcat ttctctcttc ccttanactt cttttattta ttgctatgta tctcttgctt 60
 taaagaagtt aattatgaat tgtcttttga gtaattcatg ttaacgggtgc attgttaatc 120
 cgaaaagaga gagtgatagt ttaattgagg aatagtcttt gtatcttaat tcaaccctt 180
 tctttcttaa cgttactgaa gccatttgtc aacatcctat tcttgacaac tcgcttctct 240
 aagaagacca actctctgct cttgataaat gaagccccat gaacgtctat atttttactt 300
 gaaaacacag tcatacaatg tcctttctct ttttgaa 337

<210> 30806
 <211> 480
 <212> DNA
 <213> Glycine max

 <400> 30806

ctactgatgc ggcattggcag gcttacttca ctatcttgac tccgatgcga gctatgatca 60
 ctgctcttcc ttcccgcgac gcgtcttcat atgttcgctt gagggtgctt atactctata 120
 ccatactatc cacgatgact ttggctatat caagctggca tgcttgatg gtcgttgctt 180
 agacccatat cgggttcata accgaactcc aacataactc cagccatcat tacatgctgg 240
 attggacagg caatgcttcc ccagagaatg agttcacgga tgaaattgct gacaccttca 300
 gagcactgga tagcgggttc taacgacctc tctggcggct acacttaaag catataggat 360
 gggcaacttc tcaagatgac tccctcgctt gagacaagaa cagatggcac ctcaatacaa 420
 attaaacttt cgtggagggt gaggaacaa cctcgttgat ggatcatagg cgcccaggag 480

<210> 30807
 <211> 403
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 30807

tgaagccctg acacnttgga anacaggttg ancccttgaa cccacgnatt taagacgtcc 60
cagggaaactt ttgcctttga tacggcaacc acgttgatag agacatccca aggaaaaggt 120
cgttcttcag ctaagccagg catcaatcca atctacggca ggtaatggcc catgactgcg 180
gcttggcgca taagaacatc aaggcctaca ggggtgtagta ctgatattct acgcaaccac 240
ttcccgttat ggaatccacc tttatgagat gaagcatggt gtatttacat caatctaagt 300
ctggataata aggcgggacc atctgctttt aaccaccta gaacacaaag ttgttatagc 360
taggtcgtga gagccgaaag ctacttaaac tcaagactca tcc 403

<210> 30808
<211> 500
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30808

cgccgcattg atccatcgtg agcgatgaan acttaaacc tgagaagtgc gaggacaagc 60
tctcaggcna agctcgaatg atctctgctg attctaagag aagttcacgt tcatagccat 120
cggagtctga taagagtatg atgaactatg ggacgtcatt atggtcaccg ctgaagcctt 180
ggaacgagaa accacaaagg cctcgaatga atatcactac caatgcaaag ttgtgacggg 240
ctctataggg cagctatatt gatctcaacc tccgatgagg tgataggact catcatgggt 300
caaaggcatg aacctgaagg acgaactaat agcttgcttc aagtcaaata gagaattgat 360
ctcacgataa tacgaaattg aaggattatg tggccatctt catgggtgcaa aataactatt 420
ctacacgatt acgactaagt tatagtgatt accccgattt ancaaaagg accaggagag 480
ggtcttattc ttgacctaag 500

<210> 30809
<211> 339
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30809

tatctntctt tcaattcaca cgacaataac gttntactcg gatgtctgat tgagtcccg 60
aatatatcga taagctcgaa attgaatggt gaacctctga gcaaattcaa acgacaataa 120

ctttttactc ggatgtctga ttgagtcctg tcatatatcg agacattcga aattgaatgt 180
 tgaagctctg agccaattca aatgacaata acttattact cggatgtctg attgagtcctc 240
 gtcatatatc gagacgctcg aaattgaatg gtgaacctct gagcgaattc acaccacaaa 300
 taactttttac tcggatgtct gattgagtcc catattata 339

<210> 30810
 <211> 546
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30810

cgatgtgaga cgatagancn ccenttcgttt aganccccct agctatatan gagacactac 60
 tcagaatact caagccttca acagttcaat ttcgagccg tctcgatata tgtataggcc 120
 tctaatttta catnccgagt gacaaagtag tctgtcttnt gaattgctgt caaagcttca 180
 acattcaatt tcgacgtgtc tcgatataatt acttggtactc aatctgacat gccagataat 240
 agttattgtc acttgaattg gctcagagct tcaacattcg aattctaacg tctcgatata 300
 tgaagggact caatcacaca ttcgataaat agttattggc gcttggtatg gtcagaagtt 360
 aacattcact ttcgaacgcc tcaatatatt actggactct atcagacttc cgagtagatg 420
 gtattgtcgt tgaattggct cacatgttga aattcacttc gacgcgctga tgagtctggg 480
 accaacagac tcccacaaca acttttgcgc gtacagagta gacttccaat tcattgttag 540
 cgttcg 546

<210> 30811
 <211> 562
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30811

caccacacac accgacaggg actgcgaagg gaagacaaaa acgacgagaa acgccaagca 60
 cacaaaagag aaaatgaacc ttgagacctc gaaanccagg tggaaacagc agaaccacac 120
 ggaccgctaa agacgacctg caggcaagca agcaaagatt tcagacgccg cacacggagc 180
 aggaacagcg ggaaatggac acagaagggc ccgaacagcg tagagagaca gaaagaaaca 240

agcaccceaa gagcgagaga aggaacaac caaaagagcg ggcgaggagc aaccacacaa 300
 agaaacgcac accgaggaac aaggcaagag agcagaacgc caagacntcc aaagacgaga 360
 aagacaagag gcaacatcaa gaaacgggaa agggaaccgg cgcacacgat gagaacatga 420
 aaggcaacgg gagtcgacac aacggaaagg caaacgggtg aatcgacagc aacacaagcg 480
 ccaacactag aacggaaccc gagcaggagc agcacgacaa ggcaccaagc cgcacgggga 540
 cggcagcgtg cccaagagca cg 562

<210> 30812
 <211> 504
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30812

cctgcctgtt taacacgtag tcggcgggtg cacatantcc anccctcacc gaggggatgt 60
 gacctctaaa ccccgctcagt cagttgacgc aacgggcacg accactccac agcagaagat 120
 ttttactgcc acacacatgt caccacgcct cacagtggca tatactctcc tctcgtcgca 180
 gacttgggtct acatcatatg ggtgggtgccg caaatggcag attcctggca actaaacgcc 240
 gtcgatcgac aatgactctg cgactaacct cagcatacta aggacgcgtg taaggcaaag 300
 atctgcgcta gcagacctct catgcatggc acgcccagc atgtcgggtg catctatcgc 360
 ttogaataca tcatatacag aggggatcta tgtggttagc tctgagtttg ttctcttagc 420
 gtgatgatac gtggtggcaa cgcgctcggg tcggatcgga ctcaccacta cagccacaa 480
 ctgtgctcac atagaggcgg catg 504

<210> 30813
 <211> 96
 <212> DNA
 <213> Glycine max

<400> 30813

ttcttgcttt tccttggcct acataatttt tttcagaata cacgttgta tcatgtgcta 60
 caattgtgac ccttctgatt aaatgtcaga tgatta 96

<210> 30814

<211> 514
 <212> DNA
 <213> Glycine max

<400> 30814

tcacacgttg ttgacccctt cgaaccttga tacttggcat tacgcgcctt taaacaccgc 60
 ctcactacta tttgtcaaac ctattattcg aggcgaaacag tatctatctt cccgcaaact 120
 atgtagtcca tcccctgcat ggaagtgcac ctaaaatact ataatttggg ttgccgctat 180
 atccaacaat cttcaagggtg ggcttaaggg atagataacc tacatggatt gaaccttgctc 240
 tgctacacac gaattaagag aatcatcttc ttttgactgt atggatagcg actcctcgta 300
 ttgctcttct ttccaatatc ttctgtcgcc ctcttaacga tttcatattg ctgtgttaac 360
 ataatgcctt acttcttata gggattcaaa tacccaagca ctagtgtatg cgtttgatta 420
 tctttcctct tatacaagaa acaacaacaa aggctatta attgagttgg aaaggatctt 480
 tcgcatactt tactcctata tggaacatat cgcg 514

<210> 30815
 <211> 321
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30815

tctctagggg gatccttttg ttccttcttc caaggccaag ggtaataatt ctaatcttag 60
 gtcaagggtta agggccgatg acctgaggtt ttggctcata agacttgtag agggccggac 120
 atgatgtatg taagggatat gtgttcngta accgttcagg gataacggaa tgcccatatt 180
 atttccatga tacccatgtg gacactcaaa catcangtnt gtagtaatgt gagactaagg 240
 cttangattc atttttccca tttaaataca cctagtgtnt ccagaagatg tgnnttatca 300
 attatgcatt catctgagtc t 321

<210> 30816
 <211> 319
 <212> DNA
 <213> Glycine max

<400> 30816

tgaccttgac cctgaacctg aaacgccata actcgcttga gagatacatt attcccaccc 60

cgttttttga gcagaatcct ggaaactccc gatcgatcca aaaatttagg acagtcgact 120
 attgagggga ttaaagttaa aaccgcaacc cgttgttcca aaattttaaa ttttaagtcc 180
 aggggcctcg taccctcccc ggctgagaac ccgccagggg taaaaggaca tggtgagcgc 240
 caggtttcac cgaggtcgca ttaggggggc cagactagcg cggtgaccgc gggggggccc 300
 cttaaactcg ggctccacg 319

<210> 30817
 <211> 357
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30817

agctttgcta gaaaagggtg agtattttct aananaatta tcgaaattta taatgggaag 60
 agataactta gaggcacttc ttgcccaaca aaagtgcgtt attgaaaagg ctgggttggg 120
 atacaataac aataagaaac agatagctga caaaatcttt ttcaacgcta caaaagcttc 180
 cagctcacc atcatagtat gctactactg tatgaataag ggacattctt cttttaattg 240
 ttgattaagt agtttggaaat tccaagtggg aaatacaaat gggttcctat gggaactaat 300
 aagggttgcta accaataagg acccgacata atttngtac atagatctac ctctcta 357

<210> 30818
 <211> 297
 <212> DNA
 <213> Glycine max
 <400> 30818

aaccgcccag agacagggca acacaccccc ccaggctgac tgaaccacaa ccaccaacgc 60
 caagcaccac gacgaatgcc agaagcggac cggggaaggc agcacggaaa cacggagaga 120
 agcacccgac acggcaccga gagacggacg gaaacacagg gcaggaaaca agacgacgaa 180
 caacacaccg accaaggcga acaaggcaaa ggaaccacgg aggggcccgc acacacccgg 240
 cgcacgcgcc ccagccgaa aaaacaccca aaggcagggc aagacacaca agcgaac 297

<210> 30819
 <211> 392
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30819

agcntnacat tggttatcaa ttgtctaaca taaacttctt cttttactta ggacttggga 60
tataacatgt taaagattgc agcaaagcac aacatgggca ggttgatgac atgattntga 120
aattacaata gaacattatg ccagatattt gatccatgta ccgacccac ctagtgagaa 180
aaggattggt ttgttaatgt tgtatacaaa ttcaacatac ttcaaata atagtagttg 240
tagagttcca agccaaaaga taagggtgcat ttaatgcata gtgggaattg gaattttatc 300
actaaggttt aggtggttta aatgcaaagt gaaagattgc attntactag atgaaactta 360
tttgaatgga tgcattggaa gttgtattgt tc 392

<210> 30820

<211> 416

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30820

tgctaaccce tggaagctcc taatatctcc cacactntnt gagatgggcc attcatggat 60
ggccttgatt ntctcaaggt ccacttgac cccatttcta ccaactacaa accctaagaa 120
aactatatta tctacagaaa aagtacactt ctctatattt gcatagaggg tgtttttcct 180
aaagactgaa agaacttgcc tgagatgtcc tgagtgatca tctangctcc tactgtacac 240
tanaatatca tcaaaaataaa caactacaaa tctacctatg aaatccctta agacatgatg 300
cataagcctc ataaaggtgc ttggtgcatt agtgagccca aaaggcatca ctagccattc 360
atacaaacca aacttggtct tgaaagcang tatccactta tcaccatttt tcatcc 416

<210> 30821

<211> 404

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30821

agcttgatgt gtgtattcac catctttcat agtagaatac tgggtaattgt gtctaccaca 60
cgattatcat ctccctttcca tcatttgggg gtgccactgg gctgccagat ccctccacct 120

ttgngtgtat ttctttgaag gattcatgct cctttttgca catgttctat agttgcatct 180
tatccagagc catatcagaa ttgtactgat actgcctaac aaaggcaacc gtttggtcct 240
tccaagaatg ggctcgggaa ggttccaagt tagtatacca ngtggcagct gccccagtaa 300
gactttctta gaagaaatgt atcagcagtt tctcatcttt tgtgtatgcc cctatcttcc 360
gacaatacat cttttagatg gttcttcngg caagtagtct cctt 404

<210> 30822
<211> 325
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30822

tcaacctaga ggagacggac cattccaagt gttggagaag atcaatgaca atgcctacaa 60
gattgacttg cctagtgagt ataatgtaag tgccactttc aatgtgtctg atctatctct 120
ntttgatgca gatggagggg ccttggattt gaggacaaat ccttttcaag aagcagggag 180
tgatgaggac ataaccaagg tcaaggacca tгнаacactt gaagggccca tgaccagagg 240
cagacttana caagcccaac acgtcataga gacaaagcta gtcatttgta tagctgccat 300
tgatgatgat tgaaggccca agtgg 325

<210> 30823
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30823

agcttgtttt ccnctctct tcacatgacc ctcttttctc actaagggtg gtgacccttc 60
ttcaggctgg gtgattcttc tcctctgaag ggctctatt tatagacca tcttaatgtt 120
aaaaggggaa gcaaatctct aatttacaga taatatctta ataaaattta aacaaatcct 180
aaagataaga taatatcttt ataaatataa acaataatcc tagagatntg aaattacaaa 240
tttgaattgt ttcccaacat cctctaccaa actctgttcc ttgcccctct tggcctgatt 300
caaccgctc ttccactttt ttcttctcc tatatgcttg tgtaaatggc ctattagtgg 360
ctctacattt atggacaagt ttgcttgcta ctt 393

<210> 30824
 <211> 482
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30824

taagagtnta accatgcgta gatatacctat ggtaatatct cattttttta ccatttgaat 60
 agtgtaaaaa cactacccaa tcgcctaaca aaacaataat actgtctacc acacacacat 120
 gaaaaattgn gatgttaccg tgtttgtttg acattctttg tcttcctagg agtgtatgta 180
 ataatatctt tgtagcacac atgagaccga tgttgtttgg taagagaaaa ataagattct 240
 agtaaattta gagagtttga taagcactgt gctacttcaa caaatataaa gatatgtgaa 300
 atttggtgaa gggatatctt cctcaaatac ttgtcaatta atataagcat acaaaatana 360
 attaaatata atataaatca taaaatattc tactaattat aagatcaaca ttggataata 420
 agagtaagaa cacattatta tttaaacagt tgaaagataa aagtaatatt attaaagaac 480
 ta 482

<210> 30825
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30825

agcttgtatc taacggtgat tgcaagcgat gcagcataat ctagatctct gacggaagac 60
 aagttgaacc ttaattggtg aagatagatn ggggaaatat ngaatcatga tttatatata 120
 cgtaacacgc acacttattt atattcttcc tttctctgca tctccctcta tcataggagt 180
 gttaattttt agacacttaa acaatngaaa cacttaanna taatattggc atgtgttttt 240
 tttttttttt tatctccac ttcatttaca ttacataaac aatcatatat atttcaacat 300
 ccacttattt tanacatttc atcaataact cttattnttc tctcttatca catcatataa 360
 tctatcatatc atnatnttct tcttcttttt tttcactatc tct 403

<210> 30826
 <211> 469

<212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30826

ggattcaact tgggactcct tccgttntgc ttctgacaca gcctgtctcc gctaccatga 60
 caacattcac ctccagaaca ttcttccaaa gaagaatgtg gagctcgccc ccacgatata 120
 cgacgaattc tatgggaagc tctagcggag gcaatggcat agacgaatga gaaacagatt 180
 gatgtggtgt tgggtgaagag ttctactcca acttttatga cccggaggac gactctccga 240
 agcagtgtg agtgcgngg aagaccatca aatttgacac tcagacattg aacgatntct 300
 tatggacctc gtaatcattc tggaaggggg agcaactaac tacatattcc cagtacctcc 360
 aaacttatcc tgacctcca cacctctaac cttaacatag gaccggcccg cctaataac 420
 agactcgtga tgaagatgga catggatgtg ggcagtatga tttcctttg 469

<210> 30827
 <211> 404
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30827

agcttgagta tagagacttc tcaagctatt tatcttctct ctacagagaga ctctctcatt 60
 ggattgatag gaatgaaggc tctaccctt atttatacta ctctacctcc acaatgaatg 120
 gtggagatta cttgtatcat anggtggaga ttaattctct agaatgttgc acacattcta 180
 tgagtcttta cactcttcta ctcttttcca tatecttcca taaggttcca cacatctcta 240
 gaatattcta gaggtttcca cattcttcca caagcttcta gagagtcta cactactcta 300
 gagttctcta ggacgttcta aaaaattcta tactnttcca gagatgtcta gaattttcta 360
 gaacttctcc aattaagaaa ggattccaac aattgtaatg tate 404

<210> 30828
 <211> 291
 <212> DNA
 <213> Glycine max

<400> 30828

tgagcttagt catgagaggt gtgcgtgtac ctaattctcta gagtctcatt gaagatgcct 60

cagagatgct tatcaaggaa ttactctcaa catagcttct caatgagacc gcctaggcta 120
 tgaataaaag catgtgtagc acttgtgtaa ctttgatgaa tgagagtctt gtgatacaca 180
 actcatagct cgactttctt cctttggtct tccttcatt actagctccc ccctctctct 240
 atgtctgact ttttcttttc ctccatatga acatcctctt caagcttctt a 291

<210> 30829
 <211> 385
 <212> DNA
 <213> Glycine max

<400> 30829

tgcttatctc ttggagtata taactcatta tgataagctg aacagtgtga aatctattca 60
 aacaagaatt aatgatactt gcattgacat tataaaggtc atttatactg tcactctgac 120
 acacatcgtc atatatgata actttgttga cttttgcaat aactcatctt tgaaagttat 180
 aatgatgatt tctgatttat caacaatgta aaagctttta cactaactgt acatgcctat 240
 tgagttctta ttaaaaagag cttatagaat attcatcatc tatgagcatt attaagttcc 300
 atgtaagctc tatcaaatgc ctctaaatc cttcttaca attgaagctt cgaacaaaat 360
 ggattgagac taacaataat tatct 385

<210> 30830
 <211> 180
 <212> DNA
 <213> Glycine max

<400> 30830

cctgacttac tgacagacac gcgtaaaaat tcggtttcaa aggcgtatag acgacagccc 60
 ccaaaggcac atcacaatct ataaggagat cgacataccc taccatagat catataacta 120
 tcagctagct ctttccgatc ctcccagggg gaacacatat tacatgccc aaactcaacc 180

<210> 30831
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30831

tttcttgctt ctatgcggtg gaggggtcttg gctaccccg aatgaccacc aagtgggtgtg 60
 gcatggaact ccgataagag gcatgggata aatgggttgg ccgattcaa ccatattcaa 120
 tcattgaaga gtagatagcc attatgaacg cgatactgag gtagctaga aggatcatgt 180
 tgaaccttgt cccgtaacgt ctgaatatca aagtcgtgct cgagtgtttg ttggatttcg 240
 cgcagaaagt caaactgatg aactgaaagc acgagtagct ggccggcgag agagtngcat 300
 ccgcacaaca tttgtggcac ctgcgttgta ttgtatggtg taattaagcc tagtaacttg 360
 gaaagatagt agtgcgtctc ccgggtctaa atca 394

<210> 30832
 <211> 302
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30832

ctacagctgct cgctacgaga caaaccttag aatttgtgat cagatatttc gcgcttagcg 60
 cacaaccact tacactcgct cagcgagaca tgctctntag agcacgcctt cgtaagctga 120
 gaagcctaag agcctttaat aactactaaga atagaggag ttctttatct tagtatttaa 180
 gccttggtgc ttatgaaggg ctgaacactt cattgttgat gacgtctcta ctgagcactc 240
 ttaatgtaaa actcctaact atctatttaa atgtacttgc tagtcgttca ttggctctat 300
 ct 302

<210> 30833
 <211> 386
 <212> DNA
 <213> Glycine max
 <400> 30833

tttcttggtg atccaatacc ctgatgagga tgtcccatat gttcttaaaa ctggactgat 60
 ccatttgctt ccaaagtttc atggtcttgc aagtgaagac ccacacaagc atctgaaaga 120
 attccatatt gtttgctcca ccatgaaacc accagatgtc caagaggatc acatatttct 180
 gaaggccttt cctcattctt tagaggaggt ggcaaaggac tggctatatt accttgctcc 240
 aaagtccatc acgagttggg atgacctcaa gagagtattc ttataaaaca ttttccttgc 300
 ttccaggacc acgaccatct gaaaagatat tttaggcatt acaaaactca gtggagagag 360

cctatatgaa tattgtgaga gatttta

386

<210> 30834
<211> 297
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30834

tgatactcag ctgcttctcc aaagcacagc cttctggatg attgatctgg aatgtctaag 60
tgggccagat cgctatttgc accccctatt tactaaatgc accccccttc tattattttc 120
tttgtaattc tttttccgta acgctacgag actgtgcgaa ttttgttgcg atacttattg 180
tccttgcgca gggttacgaa tccttacgga ttatgtatctt actctttttt agctttcgaa 240
gaagtactg aaactcacgg attgtgcaan aacacctctt ttcaatttcc cgcacat 297

<210> 30835
<211> 317
<212> DNA
<213> Glycine max

<400> 30835

tttcttattg ttataaagt actggatcta ttggtctagt taacttctta ccaatcatat 60
ttgttcggct aagattcttc tagatttaac tctgatccat taaatgttga tttttgtgca 120
caaattagag atgatgttga ttgtttaagg agttctcaa gtaaccaagt taaaatgggt 180
ctctttggat tgcaatctta gaagtaatgt tacattgggtt ggaatacagt gcttcaaaat 240
ttatagatag ccttcatttt ttagttaatt gagaatgcga catcctcttg gattcatatt 300
gggtgttctg tatttgc 317

<210> 30836
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30836

gcgcttatcg tctctctat ctcacgttgc tatccctttc ttcttgetca tcattgaagc 60
tccatcaaag ctacaacctt tgacacccat ttctgtcca aaatcgaga aggaagccat 120

tttcggagtc gagaagagca cctctccatt gtgggacctc acatttcacg tttgggtaga 180
 cttcttctca cataaatttt cgtgggtatt gcgttttggg agatatgatg ggtagttnta 240
 ctaggtttat gcctcatgat agttatttgt gaagaaattt gatgaaagca tgttgaactt 300
 gtcagtgttg gtatgagtca agcttaccba ttctgttgta gggttnttat gatgatgctc 360
 gtatgctgaa atggctgatg gaaaaatgat aaagatgaac ggtagaatta acctanggg 420
 taaaagtgag aatgtagtga tatgagtgga aaag 454

<210> 30837
 <211> 246
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30837

tattcatgct atatccaaga tcaaataacc cttgagggaa aaaatggtta ccaattatct 60
 aaatgacaat atgagtatca aagtccacat tctgtgtgcc ttaatttgta tgtctggggt 120
 aataattttt caaggagttt tattcttgta aaactatgtc aattcttact gaagatgctg 180
 aagctctatt attattcaaa gtacaagttc tgctagctaa gctcnaataa catatctaga 240
 gtctca 246

<210> 30838
 <211> 364
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30838

agctntattc aactgaaga ggacaaaaga gactttgttg atcaaattga gggtgggtgaa 60
 ttggaaaatt cagttgcgga ggatattcat gagtcaaata aaaggaaaac tcctttcgaa 120
 ggtttgtctt ctccatccta ccaaatttga cctgggtgtt cttagaagtt aaaattagca 180
 atttatattt attntgttat tcaatattct gattggaatt tccaaatgat ttttccaatt 240
 acagtattat tgctgatcc tttcttgaat agttgttgca cactagcttc tttgcctatn 300
 agaatttatt gatgtcataa acaacatgta ttctangtat gtttgaataa tcttctcogt 360
 aaac 364

<210> 30839
 <211> 479
 <212> DNA
 <213> Glycine max

<400> 30839

acgcacaccc tgagctgcac cacgtgtggg cgtaaacgcc tccccccccc ccagagcatg 60
 accatcgaaa accccttaag accgcccgcac agagcgcagc gtaacgagca cactccacga 120
 ttagactgct gaccaagcac cacgggagca accaagcatc gacacccac caacccgaga 180
 caataacgac ctaaaaacgc agggacaaag tcagaaaaaa atggcaaacc cgctgggaca 240
 gaaggcgcaa cacctggggg gaagaagagg gtaaacacag ccgagcaacc gacggacaaa 300
 cgcgacagag aaccgcgaga acagccgagc gattcgcggg cgagacgaag ggagcaagta 360
 cgcagaggaa accggcgacc aaaggacgaa aacaaggggc gccacacgcg gagctcacta 420
 aagacaaaag gcgagaacga cggggaaaga aaggggacag acagactcac acaccaacg 479

<210> 30840
 <211> 349
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30840

naagcttatg atggtgttca atatttatgg gggngattgt acatcaaaaa aagatgagaa 60
 aatgaaaac attntttttg taatgaaaaa tgaaaagatt aacaccaaaa agaagaaggt 120
 gttggagaca ccgatattaa tagctgcaaa gaacggtgtg acagaaatgg tagagaaaat 180
 cattgactcg ttcccagtag ctgttcatga tatggatgcc aagaaaaaaa atatagtgt 240
 attggcagta gagaacagac aaacttactt atataacttc ttgtcaaca agaaaaatct 300
 aaaggaaagt aatatattcg gaaaagtgga taacaaggga aacagtgca 349

<210> 30841
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30841

tttggataaa ctgctttgac aca

443

<210> 30844
<211> 403
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30844

agctngtgat tgttaaataat atatataaaa agaaaaattc cttgaggttt tgcacttgca 60
cgtttgagaa gaaaactcac tcgaccagga gcttgtggaa aatgccc aaa gacaattgtg 120
ataataggggt acatctgatg ttagtcactc atgcagactc cttatgattc cttatgaatc 180
caaagggtggc ctttcttgta caaattcttt cgggatcaac ccatgacatc aagtttttagc 240
aagatcaact gacccatggc atgactctat gatattaaat caggaaagtt tcacttggtc 300
acataccaaa gtgtgacaat ccattgccat ctttcaatgg ggtgcatgat cgatcccaaa 360
gccatatatt ttcttgttgt gcagaaataa tcaaagcttt aaa 403

<210> 30845
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30845

tgagatgagg aagtgttgaa gagtgaact tcctgctntt attgttgacc acagagtgg 60
acctggagat atgtcgcggg ggtcaggaga ccttgnngac gtcagggtggn gtgctattgc 120
ccanaaccaa gcttgaccaa tcccaaccca acccgggcat agtcggtcag tgagaacctg 180
tgatgtacct aagcaggcga gtcctngca gtcaacagat aanagganta caagaccaca 240
aagcaaggag gcttgtggtg gctggccagc tgtgaatttt gtgtaatatg tggattgtgg 300
tctctgggta atcgatacca naggtgagta atcgattaca aggcttanaa tngaggacag 360
gaggctaaga tggctctctg taatcgatta ccaaggggtg taatcgatta cc 412

<210> 30846
<211> 366
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30846

agcttctttg agaaatcttc tntgagaagc tagagcttag ctacacacac ncctctaata 60
actaagctca cctccttgag aaagctcctt gagaagattc ctaaagaagc tagagcttag 120
gtacacacac ccctataat agctaagctc acccccatgc caaaattcat gaaaatataa 180
aaaaaaaagc tctattacaa agactactca aaatttcctg aaatacaagg gctaaaccct 240
atactacttg aatggccaaa atacaaggcc canaagagga aaaaccaatt cttacattta 300
caaagaagaa tggatccaac cttgacccat gggctaaaaa atctacccta gggctcatgag 360
aacctt 366

<210> 30847
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30847

ntgaattcca gtccaaactc acttcacaaa atctaatttc aggtttatat aggtggcctt 60
attcgtgctc ttgcgcttag tgcacgaatg gagcgcttag cgcacgcttag tggattttgg 120
cttagcgcgc ttttctcgct taatggatga actgaagcag tgcgcttaga gagatgaagc 180
agtgcgctta gcgaacctgt acaactcatc ttcttctgga ttcttcctcg cgcttagccc 240
aggagtgttg cgcttagcgg atgctcgcta agccaacaga ttggcttagc aagaaggtga 300
aaacaacctt tttccaaagc tntcctaatt aacctanaat tgagagaaaa tgattattaa 360
acacaaaaana tgaaaatact aagtatttat tacctatact taacataaaa tacttataac 420
attacaaaat aaccataaat taagagagtn tgatgcaatn tatancaagt ttatacacia 480
aagttagt 488

<210> 30848
<211> 370
<212> DNA
<213> Glycine max

<400> 30848

agcttcagtt tagtgactat cttcaagtgt acatgaatcc taaataacat ctacttaaaa 60

gtaataaact ctttaaacca gaaaatccca tgacacaacc aagcagccgt ttgtgggaga 120
 tgagggtttgt ctcgttgata tggtccttgc ggccatttgc acatgaataa ataaaaccac 180
 agcgctgcga acccttcaca aacggcggca acagagtgat ttgcggtgcc aattttgggg 240
 tgttagggtg gagctgtgcc ttttggtgtg ggagggtggc ggtagggtgg gtgttatgaa 300
 ttttcaataa atcttataaa atatcaaag agcaccataa cacattacat tacgttagat 360
 catcaagaga 370

<210> 30849
 <211> 76
 <212> DNA
 <213> Glycine max

<400> 30849

tgaactcctt attcctttga gcataagcgg caagcttcat tcaatgtaaa gaggggcttt 60
 ccactccttg aaccct 76

<210> 30850
 <211> 391
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30850

tttctnattt ttctctcct tgatnctggc catctcttag acttcgtcct ttatctctta 60
 ttccaccctc atgttttctt gttctgcgat tcttgaaaaa tacctcattg ttgatggttc 120
 cncgactttg acgngatca tgatgtctat gccatatgta aggcggaaag tagtttcatt 180
 ggttggtgtc taagggtgaat aatgataagc ccaaagtatg ctanggagtt cctccttcca 240
 taaaccoccta gacttgtaa gtcttgtgcg cangataacc ttgtttgcta cctctgcctg 300
 attgttagtc tggggtgttc aacagaagtc acgaagtgtc tgatcaccta cctcatcaag 360
 aattcttcat aagcttaagc ttgaattga g 391

<210> 30851
 <211> 329
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30851

acctatagaa actcaagctt ggtttgaggt acttaccgt tgaagactca ttattgatga 60
agatttactg acgaacgtcg aagaacggtc aaaaaccttc gcgaaatcac ttacggaaac 120
gtttcngatg cgcctcggt cgaattctct tcacggaaac aattttacta agcacattcg 180
atagagagag aagtgcctaa ggggctgaac ccctattcta catcacttgt cccctagtc 240
atagaaaatt gtgggagaag cttgccaccc agctctccct ggcgagcagg gttgtttcct 300
ccataagcaa cagccttctg gaggaatct 329

<210> 30852
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30852

ttcttcaatg tttgcaaatt tccatgatgc cagacttttc ttctgtctcg ttcacttttg 60
tgcaataaga gtacaagacc atgtgttccc tatgttcatt tcgtcgcctt ttggacactg 120
tctcttctag aagacaacct aatggatntg ctcatttcca agcttgatag catagcctat 180
ctctcatcac caanagtttt ttaattactt gctctagctt cgtagctagg taggaagttt 240
tagttaagat catagcgtac agcttggttg acactcttat cctgcacac cacgcatata 300
taaatagtta aactgtctta ttttgatgc attaatctta tgagattctg tacataagtt 360
tcaatatgca tatatccatc actattggaa gaa 393

<210> 30853
<211> 303
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30853

nttagttaga aaatagcata gtagctnta atctaccatt atttctatat attncagact 60
gttgataat gttcgaagat aaatttcaga gagccctgag ttaccaaata ataagtgatt 120
ttttttataa aatagttaac tctattctta ccagtctgtc cccacagtgc taagatagca 180
tttgctgctt gtgacatttt aatttgattc aaaacatctt gtactgtttc aggaattctt 240

aacttanact aaacactggc taaatctgga tattgttttt gtagtcacat ctacaaacac 300
atg 303

<210> 30854
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30854

tgctntngtt taaaccccaa ttngaaatth cggagtgtcc taggtcgaag gttggaagtt 60
gaggagatta ggcacggttc caatcagaga ttaaaatact taccctgtgc ggtacttggt 120
ggagaagagg gataaaagaa acaggccaag aagatgaagg atagcacgga tagggaaaca 180
aatgggattc tatagtggaa tattntatcc acggatatta aactctatta gttcaccttc 240
tgtatgaaca tcttttttaa tagtcctctt cagagttttg aactctata aactctaatt 300
aaattgggta gattttttta gtntatcaca tacatactca attagtcaac aagttaaaaa 360
agtagaatat atatatatat atatatatat atatatatat at 402

<210> 30855
<211> 809
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30855

ggctcgacac acagacancg tngagcagca gcccgagtat ttgcanatth gcgngtcggc 60
nctancanta tcanacnnn nnncccccc cncnncccc cagcaacgag cgnncctatt 120
agtatnagcc catgcgnaag tcanncanga cancnnaann anacnacgag acangnnan 180
gcagaccgac gcgacgccg nacgacggng aangacgaca cggacacgan gacanacnc 240
acgcgctagt ataagtgagc acagtactgc gcgtcgatca acgacatcng cgaggggagc 300
gagacagacg aaggcggcga cacacatacn catcgcacga gacacagaag agtancgcga 360
cggcatctcg ccagacaaga gcatcacaga cagnggagga caacgcgngc gaagcgacac 420
aagactgcac agatgacaca gacaagagac aacacgccac gacacgacgg agcgacgaac 480
aagataggaa actcgccaca cgcagcggaa caacacacga gagagcggaa tcgtgacgga 540

ggaggcgga agaaacacgc ccacgccatc tacgaaaggg acgcgangaa caggcaacga 600
gagacgaacg acacgaaggc aggcgaacga gcaatgcacg accgggcaaa acagaacgaa 660
cgacagaacc ggcgacaacg gaacatcgcg caacgcgaca atcggcagcg cgacaacaac 720
acgaacaaac acaagcgcaa gcgatggaac aagagagggg gacccgggca gcaggcgga 780
gancaggcca gaccagacaa acctatacg 809

<210> 30856
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30856

ttgcttatat tatctctaaa aataactcta tggatgagat atttagaagt aatgatgatg 60
atgatgtata tataactctg ttattggacc caacaattgt gaagcatgtg caatgacaaa 120
attggcagac catgtgtgga acccaccatt agtaaaacaa attaactaac aagataggtc 180
tatgattaat cctaagaagc tcttttgatt atgcctttat ntgcatgaca tgagaagttg 240
gcaaaagtat attgaacata aaagttggcc gagaagcttg acttatctgt cacaagcatg 300
tcacatttct ttgagtgcag tgcactccct acgcctgctg gtgggtggaga aagacctgcc 360
cctgtcgtga tcacaaattc acgaccacca tgctcacaca t 401

<210> 30857
<211> 263
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30857

acaaatcgat gggccatctg gttgctagct tatatngcta ttaccacaag ttggcctctg 60
gttgatcggt ctctatatat tgtgtgcgcg aaaaagatta aggatatcct accaaatgga 120
ctagtcacaa aataatttct tgattgctga tattaatggg gtatttgtct gtgcagatgt 180
atgagaatat acacacatca tttcttttgc cctgcaatcg agggaaatct gcctttctag 240
gcttggtggag aacttaattc tca 263

<210> 30858
 <211> 280
 <212> DNA
 <213> Glycine max

<400> 30858

tcccatggtg catacgaatg tgtgatggat tctgtggaaa gtcacggttt cagaggtgtg 60
 gatgagagtt ttggaaactg cttctcctgt tgaatctgtc actcatttgt aagttttttg 120
 tttttcgcaa tttaattact ctgccctttg gattttcaaa tttgtggacg tgtgttgga 180
 ctcgcgtttg tctctggaag aattgttctc acaatccaac gtcgcgtgaa tgtctgttct 240
 ctcattttat tagtgcattc tacgtgttat gcctgtaatg 280

<210> 30859
 <211> 396
 <212> DNA
 <213> Glycine max

<400> 30859

ttattataac gactaataac taaacttggt atagatatatt acttattttg gatgtgtaac 60
 ggggtggagt actggctgtc atcgtcacaa gggaaatgga caaaatggca aaaaatattt 120
 tataataaag ctgtcattat aagggtttat ataattcgag aaataaattg tctctctctt 180
 taagatcgat ctatgatact atgaaggatg aaaacttcat ctttgtgaaa gacacgagat 240
 attatcgcta aaatacttat tctaaactag tagaagaatg tttataataa aaatgttcgg 300
 taagaatttc actttgataa tatgtcagag aaaatattta ttttaaaatt ctccagatat 360
 ggtctatttt gcgcaagttt cttatcatgg taaaat 396

<210> 30860
 <211> 359
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30860

agcttgtgct aatgtgataa ataaaaataa ctatnttcaa aatgaccact tttgaacagt 60
 aatttgtaat tntgcatcaa gttggtcaat cagcccttcc tttagaacgt gttttgtttt 120
 aacgtgtttg tactttgcga tgaagcagtt gaacgtgact gtaacaaatt gttgtgattc 180

tttntgttta tgtaataaag gataaaattg tttgattcac accataaacc caacacccac 240
 atattctgtt atgtgttggtg tctgctgctg ctctagagg cttcaccctt caccaaactc 300
 ttctcttttc tcttcaatca cgcacgcact tctcactcat tttccagttc acttttctga 359

<210> 30861
 <211> 124
 <212> DNA
 <213> Glycine max

<400> 30861

tattctaaat agctctcgat agcattatga atttaggatg ctgaatctag ttgactgaaa 60
 tataacctat tgagactgat gctttcaata ttttaattgtg attttttatt ataattcata 120
 atga 124

<210> 30862
 <211> 369
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30862

agctntctat atattcaaat ggtcagagct tttcacacgg aggaccgatt catgcgcgta 60
 atatatcgag atgttcgtaa ctgaacaaca gaagctctcg agaaattcaa atggtcataa 120
 cttttcactc ggatgtccaa ttcatgcgca tcacatatct agatgctcga aattcatcaa 180
 ccgaagctct atagaaatgc anatggatcat aagttttcac tcggatgtca cattcaggcg 240
 catcacatat cgagacgctc agaattgaac aatggatgct ctcgagatat tcaaattggc 300
 ataacttttc actcgcatgt gtccaattca ggggcatcac atatcgagac gctcgataga 360
 gaacaacgg 369

<210> 30863
 <211> 668
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30863

gccgagaggg cacacgggtca agacaagaca acgcnnccacc cccgcgagac tgaacntggt 60

66510-337210

gagcncntgga	aaccacacac	acaaaccaag	ctccgcaacg	agcaaaaagc	agcgagataa	120
tcctaggaca	tattactagg	cagctacact	gacgccatcg	acggggcgatg	gcggcagaca	180
agtaagcagc	gctcttgacg	cacaagagag	gcgacactag	cgaggagata	cgcattgtcgc	240
ctagtgatcg	cacgtattga	cgatagataa	tatatncgcg	cgaggcgctcc	ccacgtcgggt	300
agcgacggac	aggcgcgcac	tatcgacgcg	cgcgcacgac	gacggaacga	tctgcgactt	360
acgcgtgagt	gacgacacta	cgagacacaa	cgttcacgcg	cggcngaatac	tcgactgcga	420
cgtntagaga	ggcaagatcg	acaatatcct	ctgccctcga	gtgagacgac	tgacactcga	480
ctatctacac	acgacgcaga	cattacatgc	gaatataccg	actacaccgt	cacatgtgct	540
atgagacgca	cgagcagcga	gatgacagcc	acggcgacgc	cactaataact	acacaancga	600
cactgctgcg	cgcgaaacaca	acacaagtca	cacgaccgtg	cgctggcgaa	caaggaacgg	660
cactcgcc						668

<210>	30864
<211>	354
<212>	DNA
<213>	Glycine max

tgaatatgta caaacgcatt ggtgagattg tagatntaag cataactaga catccgaagt 120
gatcttatgg gaatggaatg gactcaattg cataagtaga gaatntacaa tgaattttga 180
tggngatgga tttttcaagt tatttggttn ttaagctgac aagtatgtca acttttgngg 240
cattaatttg tagttggtgt aaaaacacta gaaaaatcaa tgaggtgtct taagacattg 300
agaacatttc ttanattct ttgtccctaa ttctaagttt tctatttatg ggattcatcc 360
tacccaaaca accaaatttg aatttatanc aattatattt tctaaaatgt gtttagtaag 420
ggttttggtt tcgttaagag ttaatgaact g 451

<210> 30866
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30866

accatcacat gtgggactaa ggggcgggcc ggcgattgtg cacaacaagg ttttcacatt 60
cacaatgggc gcataaaccc aacaatccct tggtgccacc ttcaactgag ctcaacgtac 120
tccacgtaag ccatatcctc gtttctctca acaccgggtc cccatcaatc ctctcaagct 180
ttcacaacat ccaagcagaa caacattcaa acagcacaag ctatcacagc ccagcaaaac 240
agagcaaagg gaggaaaact cttgctcaac accaaccaaa atcacagctt tttctcgctt 300
aaaagacccc agaacaattc cttcgatcca aatcgttaac cggttgatcg actcgaaaat 360
tntaatggaa gtctctagta cataagccta catn 394

<210> 30867
<211> 475
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30867

ntgactcgga tatccgattg aggcccaata tatatcatcg ccttttatat atagaaatgt 60
actgaccacg caaatcgga cagccataac gttagactcg gattcccgat tgaagctcat 120
aatatatgga gatggtctta ggataaaaaat gaagcccatc gcanatacaa acgaccataa 180
cttttccacc ggatctccga ataagccaag taacctatcg cgatgctcaa aatttatcat 240

ggaagactcg ggtgaattcc gacgggctaa actttttact cggatgtcca attgaggccc 300
 ataatatatc atcgccctcg aatatagaaa tggactgacc acgcanattc ggacagccat 360
 aacgtttgac tcggattcct gattgaagct cataatatat ggagatgctc ttangataaa 420
 aatgaagctc atcgcanata caaacgacca taacttttcc accggatctc cgaat 475

<210> 30868
 <211> 237
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30868

agctttctact ttatgtgcan gggcggcttt cctcactttc ttgtctncaa cgcgagctct 60
 gaccactgtc cctcctttct gcggtgcttc ttttcattgc cgcttgagtg ggcttataac 120
 ctaaaccata tttcccacga tttccttggg tttttatcag gctaattatg ccgccattgt 180
 cttttgctaa acccatcccg ggttcataac cgttcccaa cataactcgg gccatca 237

<210> 30869
 <211> 480
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30869

tcaagaaaaa gatggcctca gcanattcct tatttccaga agggaattct atcaatagac 60
 ctccaatctt taatggagag ggttaccact actggaaaac ccgaatgcaa atttttattg 120
 aggcaataga tctaaatatc tgggaagcca tagaaatagg gccttatata cccaccacag 180
 tagaaagagt ttcaatagat ggtagttcat caagtgaaag cataactata gaaaaaccta 240
 gagatagatg gtctgaagag gatagaanac gactacaata caacttanaa gccaaaaaca 300
 taataacatc tgccttggga atggatgaat atttcagggt ttcaaattgt aagagtgcta 360
 aggaaatgtg ggacactctt cgattaacac atgaaggaa tacagatggt aaaagatcta 420
 ggataaatgc actaactcat gagtatgaat tatntagaat gaatgcgaat gaaaatattc 480

<210> 30870
 <211> 309
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30870

agcttcatca tttcatttcg aggggtctcg tatattacgg gactcaatcg gacatccgag 60
aaaaagttat tgtcatttgt atttgctcag agcatcaaca ttcaatttcg agcgtgtcga 120
tatattacgg gactcaatca gacatccgag taaaaagtta ttgtcgtttg aatttgctca 180
gagcttccgt attcaatttc gagcgtctcg aaatattaca tgactcaatc agacatccga 240
gtaaaaaatt attggtcggtt gaattttctc anagcttcaa cattcaattt cgagggtctc 300
gatatatta 309

<210> 30871

<211> 447

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30871

ntgagggatt tcanacgaca ataactntnt actcggatgt ctgattgagt cccgtaatat 60
atcgagacgc tctaaattga atgttgaagc tctgaccaa ttcaaacgac gataactttt 120
tactcggatg tctgattgag tcccgtaata catcgagacc ctcgaaattg attgttgaag 180
ctctcagcaa attcaaacga caataacatt ttactcggat gtctgattga gtcccgaat 240
acatcgagac gtcaaaatt gaatgttgaa gctctcagca aattcaaacg acaatagctt 300
ttttactcag atgtctgatt gagtgccgta atatatcgag acgctctana ttgaatgttg 360
aagctctgac caaattcaac cgacgataaa tttttactcg gatgtcttat tgagccccga 420
aatacatcga gacgctcgaa attgaat 447

<210> 30872

<211> 354

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30872

agcttactat atttttttat agtttgcgct atctaaaaag actttntcan aagggttgtt 60
ttggctttta taataaacia gccgagccga gctgagtctt acatagaccg agtaaaaggc 120

tcttgacaag ctgttcggct catntcacc cctatttcta atgataataa tgctcataaa 180
 aaaatatatt attaaataat atcaaaatat tcaaataaaa aatttagaat aaaaaatgat 240
 aaaggagaa aacataaacc taccgcaat cagactgtc tcaactcttc gacacctaac 300
 tttattttct attntgctat ttctagatta atcacaatga ataaatatta atta 354

<210> 30873
 <211> 381
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30873

ntgtntggag cttctatgga gaatgaagaa gaagaaagct acgtgagaga gggagaaaaa 60
 aggcttctga atttctttct tttggctgag tgaggagaga gaacagtttt ttggttttaa 120
 actaaaaggt ttttctcttt ttctattatt ttatttaagc tatgccacat gtctccattt 180
 gagtggagca aaaaggggccc actttctctt ttgattgtga cccataactca gccacaaaaa 240
 gtgagaaaaa acctaacctt tgaaacgcta aaatcttgcc tcggtttgcg tgccatttct 300
 ctggttccag ttctctgcgt ttctctgcgt ccgttggggc cagttttoga aagtaagcaa 360
 tatatatatc aaaacgatca g 381

<210> 30874
 <211> 380
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30874

tgcttcattg cctaacaggc caacttaca caggcaggct ccaagagact cagcataatg 60
 atgcatangc ccaaagttga gtatgggtgaa aagattgtat gacccaagtg aaggtgcaaa 120
 attgcaaaaa aagaatgaaa agctatacca aagcaagccc acaaagaaaa gggaaggaag 180
 tggtaccttg aacnccagt atgaatcctg ggacatttga gggcaaattg tttccaagaa 240
 ggaggtaatg atgagaatct tgaaactgac caaatacagg ctaaaggccc aagtggagaa 300
 nggatgaaag cccagtggag aaggacaaag cccccgagtg gagaaggatg aaggcctana 360
 gacagagaca ttatcaagac 380

<210> 30875
 <211> 387
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30875

tctagtcctt cagataagtg tatattncgc agttatggta tgttggtaac aacacatata 60
 ttgcacttag aattttccaa gatgtctata ttgaagtgtt taatgggtat gtactgtcag 120
 catacaagat tttgcacttg tcaaccaatt aagagtcacg tttgggtgtga ttttttgggt 180
 gggtattttc aaagtcaaca aacttactat agaagatgtc ttgtgcttgg atgatagtgt 240
 taaagcactt tacgataatg tcagagataa tgtcagagca tatacatttg tattctatat 300
 tgaatatcta ctttgctgac tcgaaacata caatctatat actcagtact tcgactagt 360
 gctgctaata taaacttttg taaacaa 387

<210> 30876
 <211> 295
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30876

agcttggatg tgacttanag caaanaagaa gggtcctgct ttccgtggac aaatggattg 60
 caaaccccgga tggtttcggt ggtaaaattt attaaaaaac ttccattttt gcaggagaag 120
 gtgaatgggtg atgggggggtt tgtacgggtt ttttaatcat tgtgcgaacc ttggagatgc 180
 tggtagattg tagaatgtac tgaaatgtgg ctacggacca aaattatcag aacagaacac 240
 aacagtcttt gtttttgttt taatttatcc atgtagagaa gtttattatg ggagt 295

<210> 30877
 <211> 332
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30877

ctttatatca tgctgttctt attacagtca tagagtcctt ttcatagtt atanttttga 60

663707 = 9074460

tcgttagata tatacggaaat ttgggggtcg atggattaaa taatttttgc ctataaaaaa 120
 attgtgtcac actacagctt agccatgttc actaccctaa acaaagtcaa tacttggtta 180
 gatcaatgta tgaacgtgta taacgcatac atgtgcatgc atgaccaca agtgcataac 240
 tgaagcccac ctttgtcctt cattgccacc aacgaagtca agtcaccaa ctttacttta 300
 cttcaaccac aagggtgtaa ttcttcggtta tt 332

<210> 30878
 <211> 355
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30878

tcgattcatt ctatgtgccc gtagtgggtcc acattgtgtt tcgtgcattt atattctcgt 60
 tttggttact ttgtataccc cctgttgacg tgcttaagcc attntactta agtcatttct 120
 cgcttaactt aaaaataaaa taaatttcca ccgaacgttt gaattgtatt atccattaac 180
 ttcggttaaa ataaattccg accgttcggt cgtgccgtaa ccacgttgga aatcaaaaag 240
 aggtaaaaaa taatacaata atcaaaaaga catcttttag taaaataaag cgganaatca 300
 agtggacgtt ttctctttgg gatntctcat tcttaatcga attgattaat aacta 355

<210> 30879
 <211> 386
 <212> DNA
 <213> Glycine max

<400> 30879

agcccggcca ccgcgagctg aacaacagac ccccccccc gagaatgagc tgaaacgcca 60
 aaacccccga agagagccac actcagacca gcgttaatcc acaccacgc ccggaaggga 120
 aaacacaacg cacaacagaa cgcacaccgc cgaaaaggaa aacaacggga gaaaagcgca 180
 caccgaccgga tagaaggaag caggcgaaca gcgaacgacc cccacaaacg gcgacggacg 240
 aaacacacgc caaaaacacg ccaaagcagc agcggagggc ggacaaggac ggacgcacag 300
 caaaggggac cacacccgac aacaagctcg aagggcggca aaaggcgcgc accgcgaaca 360
 ccgagcccga aagcagcaac tacacg 386

<210> 30880
 <211> 329
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30880

ttgcttaatt atatgtggac taaaccgggc cattggacca tgtgtatttt gcctatccta 60
 gacaacctat ataacttttt ttttatttct gcagcgcgatg caaacgaac tcaactggct 120
 gggttttcta ttcaactaaa caccttatat atactttatt ttaattcact ctntttattt 180
 ccatttatca tattcatctt ctcacaacca aacacagatt gccgtatata aatcatgtaa 240
 ataattttgg aaaaaggatg catgaatcta tgagaatgaa aatgaaaaca tggattcgat 300
 atncacaaat atgatgaaat ggtgggtgca 329

<210> 30881
 <211> 565
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30881

cgtcccacta cattccacac gccattccaa tatctgcgaa tcttcttctt cnnntaccgc 60
 ccaaccagag cgcgttgac cgtgtttgat ccttttctat aactgacctt taaatctcag 120
 cttgcgctag ggcaagtgca tctgatgatg gttgcctatg tatttgaaga cactctctag 180
 acttcaagcc attgacacca ttggctagag aatgggtgatt aaatgggagg tcaagatatc 240
 tagcgaaaga ttacgatcat tgactgagag tggctctgcat gatcagtatt tctctgatgt 300
 atctttgccca ggattatcaa gaactctatt tttcgtgctg tattaatgga acgatatatc 360
 tgtgttactc ctggaattcg aacacctagt ttcctttttt tgattgtgaa ccatacataa 420
 tccaaataag gatgctttgt ttatttgcaa actaagcaaa tatctaaata tattcttttg 480
 acaactgcgc ggtaataaga catcaaaaag attttctgct tactccactg ctgaactcac 540
 tacttttctc tgtttattga tgctc 565

<210> 30882
 <211> 377
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30882

agcttattgt attatgtaac aatgatgtca ggttttgaga aactaagggtg ttaaaagata 60
acaaggaaaa tgatagaaaa caggtaactg aacaacttgt gcgattgtgc gattgttctg 120
acagttgata cattatcatc atgggtggatt gtgactntgt gagttgtggg attccttcta 180
tatttaacat ctttgaaaca taaagcagcc actggatgtg gatgttgaat cgtgttatct 240
ctgcatcacg aattatgttg cgtccttggt gtatcgttgc ttttggcatt gaatttcaag 300
agtatgatta taaatgtcca gttcttaatc ttcccggcaa ttaaagcaat gctagcataa 360
gttttttaaa agatata 377

<210> 30883
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30883

ctgaatntac taggtttaaa natataaaga taagagggaa aaaatatcaa aaatactcac 60
aatacttgta atctttccaa tgcaaggagg tgcactctaa atactataat ttggtttgcc 120
gctatatcca acaatcttca agttgggctt aagggataga tagccaacat ggattgaacc 180
ttgtctgtag cacaggaatt aagaaaatca tttgcttttg actgtatgga tagtgactcg 240
tgtattgact tcctttccaa aatctttgtc gtctctctaa ggattntaat ttgctgtttt 300
agttaatggc ttactgctat agggtattcaa ttcccaagca ttagntnttg ctnttgtttt 360
tcttttcttc ttataataga aaaaactaan aaaggcaaaa ttaattgagt tgggaatgga 420
tcttngcaa tatttttatc actatatgga atcata 456

<210> 30884
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30884

agctngttct tcttctgngt aactacaacg atcttgggtct tcttcttctt cagcacagtg 60

ataccttgcc accttaccta ggtacagggtt tcacgactct tttgttggtt taactacaac 120
 ggtcttcgta ttgtttgttt gtttggggaa gtcgtgctca ttgcaacaat agttgtttgt 180
 ttgtgttgca aggagttggt ttgtggaact catgctcgct gcaaggaatt ggtgtgtgga 240
 actcgtggtg attgctannng gctgttgggg tgctgactat gcaaaccgtt cggggtggtc 300
 actcgtgctg gcaagggtgc attttgagtg agggtaattc aaatgttntg tgcctttga 360
 atgcttaat 369

<210> 30885
 <211> 475
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30885

tgacnccctg caanacgtga ctcgatgctc agcgacctta aagcacaact ttgctggctc 60
 aactccgagt tcttagatta taaagatata cttgtaccac tacagcttgg tatcatatca 120
 ggacaataag ccttctgaaa tcgagacaag agagtgatcg ataccgctgc tgacaagtgt 180
 aaactacacc ttgaccgtga atactatacc agaccctata ctaagctagt tactaagaga 240
 ttatatagag tattgcgcat acacagtgc tgtgagatat tatcactact catacactca 300
 attcaatcga tctatttgtc tattcacgag aactcataaa ttctcttctc tatactagaa 360
 tctcagcgag atctcctcat aatgtgcacc tacaccgaga agcaaacagt gcatcaatta 420
 ctattcatca gctatcagtg ctattttctg gtctctgagt cctaagacta attct 475

<210> 30886
 <211> 382
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30886

agcttggttg gttctccttt gccaatgtca aatctgtaac ctggtggaag ccacagaacc 60
 caatgttttc tttgtcatt ctacttatgt agtagaatat agtataccag ctccatttac 120
 aaagacttga aaagactaac cattccttcc atctttgagg taaatngacc accacattgt 180
 tgcttcagtt ttgtcaaaat acttctctca tgatcatcat tggcactctt gtcaaaaagg 240

agccttcgag caagcttctt cctacaagca atcacgcaag aacatgtntg actattcttt 300
 tgcaagcatc ttacagtang aaaaaaggta atgatgacac tcaaattagc cacagaaagc 360
 agaattcaat gatagactac ca 382

<210> 30887
 <211> 588
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30887

ggggcaggcg tgtcagacgn cgcgacgagg ataangcgac gtggcncnc ccccccccc 60
 gacgagcaac ttgaccccat ggatagaacc cantgggaan ancacacaca ctcagaanac 120
 ataacgatgc gccgcgggac cacgagacag caaggcagaa ggatagcctc gcggtacttg 180
 acaagccctc tgcgagacga gcaacggcgc tgaccgccat gggcaggagg aatcgccact 240
 gaaacgcgag gccaaagaagc tccacggccc cagccgcac actggctgac agtgagcggg 300
 ccgaatcccc tnccagcaga ggagctgcgc cgagatccca ggcgacagca cgggtggcgcg 360
 acgaagaagg gaaagaacag tcaggcctcc aaggcaaacg caaaccggg taccctgtgc 420
 cggctgaacc cgcacggagc acacgaggat gctcgcttga cgcgacacgc cccggagcac 480
 agaccggcgn ccaatagacc cagccgcccg caggaggcag acacacaatc gggacacggc 540
 agaagaccag cgttcacaca aaaacggaga cgaagagacc ggcccgcc 588

<210> 30888
 <211> 628
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30888

ctctcagcgc tctcatcgta caantcctt atactactta ctggnnttct tcaaattggt 60
 agatcgtata nngtatannn annnncaacg agtgtgcntt ttgatgcctt tgtaantcca 120
 tggcaaatac aaactcggtc ctcgtggatt ctctacactc gacctgcacg catgcacagc 180
 atgttccgat tatttgtgta tatcatcaca tgtaggtact attgaggatg tacgggcaaa 240
 cggtacataa tacaatatct cccatcatct atcaatatct acatatacaa ctccatccat 300

ccctctagtt gtccactct tcacactgaa gcacacatta cgtcacatcg tagccactta 360
 taccgcgata taactcatac acctgagccc tattctatac ctacataagc tttccaccaa 420
 tcattcaagg taatttaatc atctcactca tctcaatatt atctaaaacc aatcaactat 480
 agtgaacata gcacaatata ttgaaccata cttactccta aataataatc ttctacaaca 540
 tatatgaaac cctcataaac aattacgtat atcatacggt tcatcattct aagaatttat 600
 atggatatcc ctaatcatat gccataat 628

<210> 30889
 <211> 337
 <212> DNA
 <213> Glycine max

<400> 30889
 agcttggtga atgactggac atgatatatg tcagggtggt gggtcggaca gcagtccagg 60
 ggtaaaggga tgtcccatat tatttccatg acacgcgcgc aacaatgatg attcagaaat 120
 tctatgcaaa actggtcaca cattcaccta tgtggacact caagcatcaa gatttggtgg 180
 tcacgcgaca ctatggctca ggattcatta tttttcctat ttaagtcaac tcattgtttc 240
 caaaatatgc tccttgatca aatcatgcat tcatctgagt ccattttggg cgttctggaa 300
 aattatctca acattcaccc ttcagggtgcg tacatat 337

<210> 30890
 <211> 299
 <212> DNA
 <213> Glycine max

<400> 30890
 atgtggaagt cggcctatgg atcactacat agattacaga gacactaatc acgcataaac 60
 cctccatagc atgatgtcct aattcaactg aactcactta ctacaacgaa gcccatatgc 120
 tegattctct caacactcgg ggccgaatcc atcctgcaa gctgtaccaa cctccgcgta 180
 ctacaacatt caaacagcac aaaataggca tccaggcata acaaggcaaa ggcggaacac 240
 tctgccctaa acaccaacct agatcacagc ttttatgact gaaagacctc agtaacaat 299

<210> 30891
 <211> 320
 <212> DNA

<213> Glycine max

<400> 30891

tacatatgta ctgacctaga gccaacataa cactataggc atgcatcatt accgttgaca 60
ctgcagccga tgcgattact atatgctaca tgctccatat tgataccaat ctacatggca 120
caccttcaat acgtacactt ggcaaacaca ttcaactctt caacctcatg ctcacgcttc 180
tctttcacat cttcacctaa tggcctacga tgaccttcat ttgcatgtac tgtctgacta 240
gcgteccgtc ttatcttcga cagcatcatg catatcctct tcatgcgcca ccgacgaaga 300
aaactgtaat tatctatcgc 320

<210> 30892

<211> 97

<212> DNA

<213> Glycine max

<400> 30892

agctgtttga tatattatgc tcttgaatcg gacctcctag ttctaagtca tgaccattta 60
actctcctga tagcctccgc agatcaatct tgagcct 97

<210> 30893

<211> 412

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30893

ctatagcaaa ctcaagctnt caagaaattc aaatggctct aacttttaac tcggagggtt 60
gattgatgtt tataatatat cgacacgctc caaattgaac aatggaagct cttgagcaat 120
tcaaattggtc ataaatagtc actcggaggt ccgattcatg cgcataattt atcaagacgc 180
tcgaaattga acaacagaag ctttcaagaa attcaaatgg tcataacttt taagtcggat 240
gtccgattca ggcacataat atatcgagac tcacgaaatt gaacaacgga agctctcgag 300
aaattcaaat ggtcaaaaact tttaactcgg atgtccgatt caagcacata atatatcgag 360
acgcgcataa ttgaacaacg gaagctctcg agaaattcaa atgggtctaac tt 412

<210> 30894

<211> 240

<212> DNA
<213> Glycine max

<400> 30894

accatagtca tggataggaa ccggaagaag acgcccctaa tatctgtcac ctgtatcaga 60
tatcatggtc agaagctttt cgccaaactc agcaatatga ccaatgtata tgcggacata 120
gcaaaatgct cccagatgct caaagagtga gcatctggcg ttgaatattt ttctattctt 180
cttagcacia ttggaatctc tatgagcctg atactatcac cttatctcat aaaagatcct 240

<210> 30895
<211> 200
<212> DNA
<213> Glycine max

<400> 30895

agcttgactt gagtcatgaa gagaatataa atatgtggcc atgttatgag gtttatataa 60
tcacctctcc aacgatctta tcaactatca atcattcttt ggatcatcct atctttcaat 120
tcttttttaa catccattgt caaacatttt tcaatgaatc tttcaatagt ctttctatgg 180
aaattttcga ttcatttctc 200

<210> 30896
<211> 259
<212> DNA
<213> Glycine max

<400> 30896

accctttatt gctaattcat ctctagaac tcaagtacaa agtctacctg acaatctttc 60
attctacttt acttggcata atacattaca ggacactaaa cccaagtacc ttgtaccaac 120
ccaagctgct tgtacaacct atataccagg ccccttgatc attctaaaac caagatccct 180
tatcgtactc tacactacac accatgcaag tcaagtaaca aacaactccc caactttgca 240
cactcatcag gatgcatac 259

<210> 30897
<211> 574
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 30897

cacacccact cacgtcagta actnagnaana atatantaat aatcgaatac tccccaaact 60
catgtncagg ataattatnt ccacagggct acaggnacnag gttgaaacca tgtagaacc 120
nnctgcaann acgcgacacn anagaanaact caagctggct tgccttaaga tgacaaaggg 180
atgttataaa aattctatcc atacacactg agtcctcaag aaaggctgca gacgaccatg 240
atggctttcg ggtgtcaaag agtctactta gactctggct gaataaaggg agactaaagt 300
agtctcggac ggtgcacatc gagtcttcga caaaagggac agacgaccat ctttgtctct 360
gcgtgaatca cacttgattg cctcaggatg acgaggggga gacctaaga cccccagtcg 420
ataaacaacg agtccacgaa caacagtgc caccaccatg tcgggtctcta ctcttcgaca 480
aacatgattg cctccggatg aagacgcgga cactaacgta atctcgaacg aacaacatct 540
acacctcaac aaataggcgc acaaaaccat gtcg 574

<210> 30898

<211> 379

<212> DNA

<213> Glycine max

<400> 30898

ctgcgcgctt caatcttggt gtaaactgaa taggccacaa cggtcgagac cacacatcta 60
ttcttagctt aattaaactg cagaatcaaa tcataagcac aaccagttct gtataactat 120
caatcacaac ggccatcaag tgataaccaa ctaataaaca agcgaaagta catacgatga 180
tggtcaacgg attgtggaat tgaaactcaa gaattgaaca accttcgaca taatatcagt 240
gcttattata gaaactgtgg aggtatcaga atatgtgtga aagaaaataa cacttaatat 300
taactagatt aatgctgact gaatactatc taatgtcttg acaagtactt cgtgataata 360
cctggaatat ataccttca 379

<210> 30899

<211> 414

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30899

tggggattta ttntagga aaactcagat acagacattc tttttcatag gtatttttgg 60

tgttcttcaa ttagaattgg agttttacct atgtaatata tgttgatctt ttatagaaga 120
 ttttattacg tggattatca agatgaaact ccaattctga tcggagaaca aactaaaaa 180
 cacttaagaa actacaccta agttttgtcc tttattttat agtaactttg ttcataaagt 240
 tactagaatg atcaataaac tacaaaattg tgggtgaata ggaactgaga cgtttcccaa 300
 ctctccaagc cgaaatagca gcagctacaa atgaggcttt agagagggtc cgcgaaagaga 360
 gtaagaagac agctatgcgg cttgtggaca tggaagcttc ctatctcact gtgg 414

<210> 30900
 <211> 397
 <212> DNA
 <213> Glycine max

<400> 30900
 cagttttcca cttgtactcg tgataaaatt aaagaaacat gttagattaa gtatcccaca 60
 atttaagcaa gaataacttc attttggctt ccaaccttac tggatatctag gcattactat 120
 gtaacacata cgttttctac tatgacatag tgcataagat ctaccaataa ttttcaagta 180
 ctaattaatt aaataattga aagttgaaac tacactatcg atatacattg attagcttca 240
 caacttgcta aactagaac actgaaacat tcttcatttt acacaaaaaa tactaataag 300
 aaataaaaag actgcgtggt gttggaagaa ccaaaacgta gcaaactaca ctataactca 360
 ttgcttaaag catgaacaat cttaacctga ggaaaaac 397

<210> 30901
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30901

ntgcggattt ggtcttcgcc agagaattga tcgatgtggt ttctaacaga ggcaaatttg 60
 atcatcctac taggacgact gagaaaactg gggcaaataga agagggtgag aaagagggag 120
 aaacccatgc tgtgactgcc attcctatac ggccaagggt cccaccaaac ccaacaatgt 180
 cattacttag tcaataacaa acctactcct taccaccac ccagttatcc acaaaggcca 240
 tccttaaate aaccacaaaa cctgtctacc gcacttccaa tgacgaagac cacctttatc 300

<210> 30902
 <211> 383
 <212> DNA
 <213> Glycine max

<400> 30902

agcttatgtg attatagtag ctcagctgta tcattgggta aagaattggg gggcattgaa 60
 cttttggctc ataggttaca gaaagaggta cacagagtca ttggtttggt tggaggaact 120
 gataacatga tgcttactgg tgaaagcttg ggacatagta ctgatcaatt gtactcccag 180
 aagagactca taaaggtctc ccttaaggcg cttggttctg caacatatcg cacctgcaaa 240
 ctctaccaga tctcaacatt ctcaagacag ttcattacct ataactctaa acttgatttt 300
 taaaaatgaa gataaagttg gaggtgacat ttattattca gctgtactga tatgagttaa 360
 taattcaca aatcctacct ttt 383

<210> 30903
 <211> 475
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30903

cctactccac caatattatc tttatataa gtgtatttat nctatattac anctctacna 60
 caacagaaaa ttgagcatga anctttgaac cagcagaccg ggatcttgga gcaccagcag 120
 cagcagcgtt ttctcattcc tctgtttacg agcaaaagta gacagcctac tcgttagact 180
 aattaaaact aagattccta ctctatccta tgctggacta gaccagctta taagctgaca 240
 aagttagacc aattagccta agcatagcct cattcccgtt attggactag atgagaccaa 300
 caacattatt ctaacagcat atcttaaacc aaacttaatc cgcaaccctc attaagacta 360
 gattcatcct gctaattaaa gttaatgcac agaaatttcc atgctaagta cctagcctgc 420
 cacatagggg gacgaccaca gctacaaatc tatcacttaa tgagcatgac acacg 475

<210> 30904
 <211> 305
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 30904

agcttatgga ttatagtagc tcagctgtat cattgtttaa agaattgggg ggcattgaac 60
ttttgggtca gaggttacag aaagaggtac acagagtcac tggtttggtt ggaggaactg 120
ataacatgat gcttactggt gaaagcttgn gacatagtag tgatcaattg tactcccaga 180
agagactcat aaaggtctcc cttaaggcgc ttggttctgc aacatacgca cctgcaaact 240
ctaccagatc tcaacattct caagacagtt cattacctat aactctaagc ttgattttta 300
agaat 305

<210> 30905
<211> 411
<212> DNA
<213> Glycine max

<400> 30905

tgtaagtatt cgtgactcat gagattcatg gaaagtaagc atttgttctt actcaagtat 60
gtagaattat taatttttgt tgttatcctg actctggaga tctgcagttt aaccatatgt 120
tttatggaat tgttcaatta caacatagtt cacttttttag taaatattaa aatgaaagtc 180
tacttgtttt acagagataa atagatgttt tgcacgggtg aattttgtta attagtgtga 240
gcttgaagat gtatgctaaa atgactctgt tactagttaa atttggcaaa aaaataaaaa 300
taaaaatcag cattctacat ctacatcggt tgtagaccaa aaacgatgta gaaactctac 360
attctacatc ggttggatct cataacgatg tagaaacttc acaattctac a 411

<210> 30906
<211> 93
<212> DNA
<213> Glycine max

<400> 30906

tagcttttta ctctatacta agtaatgagc ggtcacttct gagacagata tatatacata 60
tatacatata tatatatata tatctatata tat 93

<210> 30907
<211> 400
<212> DNA

<213> Glycine max

<400> 30907

agcttcatcc tcagatccct cttgttggac taggcttaat ttagacagcc ctccctagggt 60
tagactaatt taaactaagt ttcgtcctca gatccctcat gttggactag acacagctta 120
aatagcttac aaaagtttag actaatttag cctaagcttt gtcctcatat ccctcttatt 180
ggactagact tagaccaaac aacattattc taacagcata tttaaaacca aaacttaatc 240
cgcagatccc tcatttaaga ctaagtttca atcctgcttc attcaagttc taaggcaaca 300
gtacatttcc caatgctaaa gtcacctaac catgcacaca aatgggtgat cagacaaaaa 360
gcatacagaa ttttaagcact aagagaagca ttgaacacaa 400

<210> 30908

<211> 372

<212> DNA

<213> Glycine max

<400> 30908

tactgcgaca tgaaagagtg gtcagcgctt tcagatttct cctagcccag caggctgtca 60
tgaggactct ccacatgccc acacacgata atctttcata accttcttcc gacaatgac 120
tctcatgcag ctcatactct taagactgtc actctatatg acatgctcca tattatatgc 180
agcatgcagg tcatatcatc accatatact atatggactg ctacttctgt cgacaagatc 240
agtctcttta gaatctataa ctgatccaga cccaaccgac tggagtata cctatgtaag 300
agcattcctt caaacgtttc tttgaatctt caatacgcag ctacatatc tcaaacta 360
gcaccgtaca tg 372

<210> 30909

<211> 402

<212> DNA

<213> Glycine max

<400> 30909

agcttcttat ccaaggetca tcttgggtgt gaagctcctt ctccatggc ttattcctta 60
atggatggcg cctcctctca cctccttcc tttgtcttcc gctgcatctc catggtggaa 120
aatcaccatt aaaggacccc attgaagctc aaagatccag cctccataga agccccacaa 180

gcaagcttcc atcatagaag ggtcagaatg gtcgaggcag atcaaagtca agaggggaaag 240
 gtcaaagaaa atttcaaagt ggcattactt gttggaattg tgacaagaga ggtcacttca 300
 gcaatcagtg taagccacca aagaagatca agtcgcacaa aaacaagaag cgcgatgatg 360
 atgaatccac aaatgcagca actgatgaac ttgatgatgc at 402

<210> 30910
 <211> 416
 <212> DNA
 <213> Glycine max

<400> 30910

gcgcgggtct gggagacgaa gggcaagtgg aacgttatat acgattatga tgttccgagt 60
 acattggatt tggtagcacc atgccctcct gatttccagg tgggaaattg gcgagaggag 120
 gaacgccttg acattgactc agcgagcata atgtcaacct ttacggatct aaaagctcta 180
 tagctggggc taggctttag aagttttcct ttgggtaagg ctttgtgact ttcgtttttg 240
 aatttataat acaaggacct tgtttcatct gttcctacgt atctacccat tctcattcat 300
 ttgcatgatg acttcttttt ctaaaacggc agatccgatg acgagtcctt cgaagggtatt 360
 aatacctggg acccgcttat caacttcgag caagagatga atcacaccga atatga 416

<210> 30911
 <211> 349
 <212> DNA
 <213> Glycine max

<400> 30911

agcttgtaat atgtctagcc aactatatgt tcagttactg gtggcctgtg gaagatgatt 60
 gggtttttta ctgcattgta taatgaatga tcgaggccgt acccgaatca aataatcatt 120
 aaaaatacag tatttaggaa gtgacccatg gtcgtctccc aacgagcaat ggtcaaccaa 180
 atgttcataa cagatagtaa taaaacagta acgaattggg gggggggggg tgtttgttta 240
 tagaaactac acaatcataa aattctaatt gtactatatc agagataaat catgtagtat 300
 caccttgatt cacaagctag gttcttatcc tatgatacca tgattttatt 349

<210> 30912
 <211> 428
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30912

ctcaagcttg tggagcttag agaccttgta naaagtcacg tgggaatatg gtttattcta 60
tacttgctg atctgaagag gttgtctgag ttctgggtcaa gttactgtga gcaaagaaca 120
tactactact gttctgtggt gcatatagtg atggagctta taaaattgat tagtggatat 180
ggggagaatg gtagattgtg gcttatgggg attgcaactt ttggaacata catgaatggt 240
caagcatgta accctccct tgagactatt tgggatcttc acctctttac agttccagtg 300
ttactatcat ccttgagact agcttaactt ttgataggag ataattatcc ttctatatac 360
ttcttattta tgcttgagtg tacaatattg atttcttagt gatattaata cgctgggtgtt 420
tttgtctc 428

<210> 30913

<211> 133

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30913

tctacgctta ccgcaggagg agacatacaa cgaggagaa actccacgtt gtacgtcgcg 60
cgcagctcgt caacttgag gaactatacc gtgttacatg cgctgctcat gaagacgatg 120
tataaaccca ctn 133

<210> 30914

<211> 319

<212> DNA

<213> Glycine max

<400> 30914

gtaatgaacg atatacctgt gagggacagg gccgatacgt taaccactat gaaagtggaa 60
acaattgact tcccatgcga tctatcaagg acacgcgttg taataagacc accgacgagt 120
acctttggta gagaatccca caccaacctt taaaagacga atcccatgg tgaaagtcaa 180
tgaagaatta ttgctcacag tttttgggtg accacacgag atcggtacta gaggctcgct 240
ggctagactg ttctttcttg cgaatgtgga ggagcatttg cggatgtgat taccataacg 300

ctaacttgaa tatccccga

319

<210> 30915
<211> 413
<212> DNA
<213> Glycine max

<400> 30915

tcatgatgaa tcaagatcgg ttcagagatg ttctgatgat atcaaagatg aagaccaagg 60
tgatgacgaa aagctcagcg ctcaatcata gaatgagttc aagatgggtca agatagaatc 120
acgatcactt caagactcac gaggaaagtg gaagaactct tcgagattca agaggaaagg 180
tgagtcttag aatcaagaat cacgattcaa ggatcaagct ttcgagaatc aggatcaaga 240
ttcaagactt aagactcatg aatcatgaga aggcttaatc aatatcagta tgaaaagggtt 300
tcttcaaaaa ctaagtagca catggatggt tctccgaaca tgtttaccac agagtgttta 360
ctctctggtg actgatcacc agactgctgg aatcgattac cagtagcaga atg 413

<210> 30916
<211> 387
<212> DNA
<213> Glycine max

<400> 30916

agcttctatc caaatggact taccttgaat taattcctta gatagccctt ttgagccttg 60
tttctttttc cttgtttaga agctcactac aatccttatg tgaaaaacca tgatattacc 120
atataccttaa ggaatttttg agcttttgaa ttgttttggg aataacagtg gggggttttt 180
gtttcattga acaacttgta ttgttggtta tgcttcatga tgtatgttgg gccatacttg 240
atgtacattg tatatgggat aaatgatgga catgctgaat gaaatgttgt ttctcaaagg 300
ctatacagta aaaaaaaaaa taaaattcga aaaaaaaaaa cgaataatag aaagagatca 360
gcaataaagt tgagtgaata agatctt 387

<210> 30917
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30917

tgtatacatg aatttgattn ttatgatgct canaacctat atgttgggaa acataaactt 60
catcttcagt gatccattta gggaaacact cttgacatcc atttgggtcta actntaaatc 120
cataacataa gcataagtag taatcttacc acttctagtc tagctatcgg tgcataagct 180
taaccaaaga ctatactggt ttgttgggta tagctcttga ctactatcct tgccttattc 240
ctagtgatca aaccatgttc attcaattta tttttaaaca ctcathtagt gtaaagtatg 300
ttcatgtttt taaaataagg tattaattcc catatatcat ttcttttaaa ttgggttcaac 360
tcctcatgca tggacatcat ccagaactta catttgagt cctcttctat agacaatgg 419

<210> 30918
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30918

agcttgagtg cttntgctgc aggacaagct gcagccttta agatgtttga aacaattaaa 60
aggaagccag aaattgatgc ttatgacact actggtcggc agcttgatga catccgtgga 120
gatatagaac ttagggaggt ttgcttagt tatectacta gacctgatga actgatattc 180
aatggattnt ctctttcaat accaagcggc actacaacag ctttggtagg agaaagtggg 240
agtgggaaat ccacagttgt tggtttgata gagagatttt atgatccaca ggcaggtgaa 300
gttctcattg acagtatcaa cctcaaagaa ttcaaactga aatggatcag acagaanata 360
ggcctagtta gccaggaacc agttctcttt a 391

<210> 30919
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30919

gtgattggac atggttgaaa ggataattat caacacaatt atacgggtttt ttgtagatac 60
tactcaaaca ttggttacct cttttagtn tagctcattg gggggcaaata ttaagccact 120
cattaagatt atttttattg acaacatgcc tgtgtgtaaa atgatcaatt aggatttgac 180
agtgtcaaac aatgggtntg gatttgtatt cagctcaatc tatgcaagaa aaacaaaaaa 240

ggggatgagc tctgatacat gcagaacata attaatcttg tcacctgctg gaaactggct 300
 aaatccctgt ttatatcttg ctgttgaatt tcatggatat gctgcttgaa gatttgatta 360
 gattataatt tatacatatg gtatgacttg tga 393

<210> 30920
 <211> 374
 <212> DNA
 <213> Glycine max

<400> 30920

agcttttaag tgcgggttcg ggagacaaag gtcaagcgtt cgcgatatgc gaagatgata 60
 ttccgagtac tttggatttg gtacgaccat gctctcctga tttccagctg ggaaattggc 120
 gagtggagga acgccccggc atttacgcaa caagcataat gtaaaccctt acgggtttta 180
 aaagctctat agttgggcct aggctttaga gttttcattt tgttaaggct ttgtgtcttt 240
 tgtttttgaa ttataatac aaggatcttt ctccatctgt tccctggtctc taccattct 300
 cattcatttg catgtttact tctttttcta aaacggcaga ttcgatgacg agtcccccca 360
 atgtactaat acct 374

<210> 30921
 <211> 402
 <212> DNA
 <213> Glycine max

<400> 30921

tcaccggatg atgcagatcg aacatttctt aatctatatc atccaattgg tattcagcga 60
 ttgaatagaa taaacaatgg ccggtgtcgg tcgttatatg gccccgactg atatctttca 120
 gccgacattg cgcaatttct tttaaaaacg cttgccgata atgttttttt tttttttacg 180
 gtagaggaag tttttggttt tgggtgttgc taaaaattt acaacgtaag tcggctaggt 240
 ttttccgtgc gagctcaacc gagggttcgc tcccacagac actggcatgt tgttcttctc 300
 atttatgagg acaagataac gttggcccat cccggcaaaa acaataaaa aacattattc 360
 accgaaattg atcgaaaaaa atgatagctg acgtcggaat gg 402

<210> 30922
 <211> 384

<212> DNA
<213> Glycine max

<400> 30922

agcttattct tgtctgaggc atcttccaag tcaatatctt tttcttcctt gagttcatca 60
tatatccaaa acggaaagta atcattgctt gaaagttgat ctggaagagg atctgagttc 120
ctccttctac ttgccatttc catcaaaagc tttccaaaac tataaacgtc ggccttatat 180
gatactccac caatattttt gtagtataat tctggagcta tgtagcccaa agttccaatt 240
gcttcaggta aaacaagaga cctatctttc acaggatgta gctttgcaag tccaaaatct 300
gaaacctttg ggatgaagct ctcatctaga agaatatgt gtggcttgat atcaaaatgt 360
agaatttgca catcacaacc ttca 384

<210> 30923
<211> 328
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30923

tcccgatatcc gtacttggaa ggatctgatt actgcctttc taangcaata tcagtataac 60
tccgatatgg ctcccgatcg cactcagctg cagaatatgt tcaagaaaga gggtgaaacc 120
tttaaagaat acgcacaacg gtggagagac ctggccgcac aagtggctcc tcccatgggt 180
gagagagaga tgatcaccat gatggtagac actctgccag tgttctacta tgagaagcta 240
gtaggttaca tgccgtccag cttcgcgac ctagtgttcg ccgggggaaag aatcgaggta 300
ggattgaaaa gaggaaagtt cgattatg 328

<210> 30924
<211> 379
<212> DNA
<213> Glycine max

<400> 30924

agcttgagct atcagaagac ttgcttattc atttagtggt gattttctcta ccttcacagt 60
ttagtcagtt taagatctct tataactggt agaaggagaa atgggtctctt aatgagctca 120
tttcataccg tgtgcaagaa aaggaaaggc tgaagcaaga aaggactgaa agtgctcatg 180

ttgtgagtag ctctaaagac aaaggcaaaa gaaaaaggac tgaggagccc aagaatgaag 240
 ttgttaaggg tccaagacaa aagaaacaaa atcagggtga caactttttc tcttacagta 300
 agcgtggaca tgtagagaag aaatgtaaca aatatcatgc ttggcgtgca aagaagggtta 360
 tgtttcttac tctagtcta 379

<210> 30925
 <211> 416
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30925

ntacaacacg gccaaagtga tcttntttgt ggggaacgat gctctggtct tccaccatga 60
 cttgtccata atcattccgg caacggagga aatcagaact tgtgagaggt tattaatgac 120
 tntataaccag aaccatctga aacaaattaa gaagatcgaa caaaataaaa gtgaattaat 180
 attgcacatc tatactaaat caaatgatgtt gaacaaaggc ttaactaact aattatgagt 240
 ataccttgta gctgcatctt cttgaagatt ttagaatcga tcacagatat gattgggtggt 300
 tacagcgtga atggaagaag actgaaaatg cctatgatgg agtagctatc caatacttga 360
 tcattctatt gggtgtatca acttctcggt ggtatataca ggaatggtct tggatc 416

<210> 30926
 <211> 402
 <212> DNA
 <213> Glycine max
 <400> 30926

agcttgaaga ggatgcttta atggaggaaa agaaagaggg agagaagtgg aactttgaag 60
 tgtatctcat aagactttca ttcatacaag ttacaacaag tgttacacat gcttctatct 120
 atagactagg tagcttcctt gagaagcttt cttagaataa cttccttgag aagcttcggt 180
 gagatgctag agcttatcta cacacacca tctaataact aagctcacct ccttgagaag 240
 ctagagctta gctacacaca cccctctaata aactaagctc acctccttaa gaagagaagc 300
 tagagcttag ctacacaccc ctataatagc taagctcacc cccatgacaa aatacatgag 360
 aatacaaaaa aaaaatccta ctacaaagac tactcaaaat gc 402

<210> 30927
 <211> 407
 <212> DNA
 <213> Glycine max

<400> 30927

tacggaccta tgaaactcag cttttatcca ggctcatctt ggtggtgaag ctccttcttt 60
 ctggcttatt tcctagtggga tgacgcctcc tctcacctct tctcctttgt cttctgctgc 120
 atctccatgg tggaaaataa acattaaagg acctcattga agctcaaaga tccagcctcc 180
 atagaatccc cacaagcaag cttccaccac aagtagtata aaacggtaag aaccgagtat 240
 cgaactctcg gggaaacttg gttatctggc aagctatttc gataaataag cgtctgggtat 300
 ggaaatataa ctgtgggttat gaacagggtat ttaaaactatc taggcaaaaa gaaagaaaat 360
 cacgtaagag aaatactatg taaaaacaag tagagaaagc gttggtc 407

<210> 30928
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30928

agctntgcgg atttgggtctt cgccggcgaa atgatcgaag tgggtctaaa aagaggcaaa 60
 tctgatcatc ttgctttgat aaatgcaaaa aaagactggg gcaaataag agggtgagga 120
 tgaaggagaa cctcgtgttg tgactgcat tcctatacaa ccaagtttcc caccaacca 180
 acaatgtcat tactcagcca ataacaaacc ttctccttac ccaccgcca gttatccaca 240
 aaggccatcc ctaaaattaa ccacaaagcc tacctaccgc acatccaatg acaaacacca 300
 ccttttagcat aaaccaaacc accaaccag aatgaattt tgtagcgaga aagcctgtag 360
 aattcacccc aattccagtg tcctatgttg acttgctccc ata 403

<210> 30929
 <211> 426
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30929

ntanagactc aatttcattt gaatgtgggt gggacatata ttgagttggt tattaaatca 60

atatttattt ctaatttgga aaaatttagt tattcttggt caatttgaaa taataaagtt 120
 agtctctctc gtgattcaaa tggtgtaagt tatgaattct tttatggata ttgagtgttc 180
 ctattggaaa caattttaag tggtaaaaa cttaaaatta aatgagaatt tcatcacttt 240
 ataacataag tgctaagtca tatctcttaa tagagaattt ggatgcttgt gttataagaa 300
 attttggatc ctttatattg gttggacctt atagtctaca ttgagtatat aaaggcacia 360
 taaacaaaac aaaaaaatgg aattcatgtg ctgaaagaaa taaggatgtc ctggaactaa 420
 tacata 426

<210> 30930
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30930

agctttgttg gatttctgcc atttggctct attttgatac ctatccatat aaaataataa 60
 atattttagt caaaaaata tatacaagtg ctttacaaga tcgtagtgc aacatttgag 120
 ttccatgtcc ttatttctaa aagtggatg aagaattttc cttgagaaat cgaccacccc 180
 atgtccttat caccttgggt aaataaattt ccttttgcaa agttcttatt tctaaacttt 240
 attctacatt ccatgtgatg aagaatttga cagtctgttt cttcggatgg ggaccatttc 300
 acaaaactct aagctttcca gaagaatata gaaaaccacc gcataatttg tttcataat 360
 tatacgacct tatgatatta gtcaatgact atntaatat 399

<210> 30931
 <211> 390
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30931

tcatgatgat gaatcaagtt gattcaagta gttttaatga tgaanaagat tatgacaaaa 60
 agcctaaaga atgatttcaa gattaagttc aagatcaaga ttaatttcaa gattcatcaa 120
 gaagattcaa gattcaagaa taatcaagat caagattcaa gactcaaaga ttcaagaatc 180
 aagagaagac ttaatcaaga taagtattaa aaagtttttc aaaacattga gtagcacaag 240

aagatttcac aaaattatta ccaaagagtt ttactctctg gtaattgatt acaagaatgt 300
 agtaatcgat taccaatggt nttacaacgt taagatnttc aaaattcaga atgaagactc 360
 acatctgttg atgtgtaatc gattacacct 390

<210> 30932
 <211> 384
 <212> DNA
 <213> Glycine max

<400> 30932

agcttctctt ggaccttgaa caagcaatca actcctcttt cagaaccctg ctatgtgctc 60
 gcgactgggc cctttcttcc cttegcaact tgagttcatt attgctaccc catagagctc 120
 cgcgaaatth gttccggcca tactcttctt tgcgagccct cttgggtctct tgttcaaggg 180
 ctcttgcggt aattgcattc tcttcccgta acccggcaca ctcttccga acgtgtgtag 240
 cagccaactt gaacttctcc ttggcgagtt ttgcctttcc taactcgctt ttgagagctt 300
 ggacttcttc gtcctcttcc ggtgcttcaa aattctcttc gctgacgact ctttaacttg 360
 cgagccaatc taaacctcgc atgc 384

<210> 30933
 <211> 340
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30933

gcntnnanca aatcatataa gataaatgca ttcattgcaat ctgtagatat atcctcccaa 60
 acgtcaaatt ctccgcctat atattcaacc tttccatcac tggcacgtgg agtgaatctt 120
 tctccatggt gcaatactaa agttatattg tcattcattc tacacaatta gaaaccgcaa 180
 acatgggtcag atattangaa ataaaaaac ctacctcaaa aagcgcggaag acattgacat 240
 tgtcaaaaac cgcggaagaca caatcaataa ccaaaaacat tgtcatctat aaaaacagag 300
 catcataaac gaacatatta accgatcata aacctcccta 340

<210> 30934
 <211> 396
 <212> DNA

<213> Glycine max

<400> 30934

agcttggttaa tccatggaag ctctaatat ctcccacact ttttggggtg ggtcattctt 60
ggatggcctt gattttctca ggttccactt ggaacccatt tctaccaact acaaaccta 120
agaaaactat attatctaca caaaaggtag acttctctat atttgcatag aggggtgtttt 180
tcctaaggat tgaaagaact tgctgagat gtcctaagtg atcatctagg ctctactgt 240
aactaaaat atcatcaaaa taaacaacta caaatctacc tatgaaatcc cttagacat 300
aatgcataag cctcataaag gtgcttggtg cattagttag cccaaaaggc atcactagcc 360
attcatacaa accaaacttg gtcttgaaag cgttttt 396

<210> 30935

<211> 390

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30935

ctggggaata atatttaaca actggagtag tgaacttaac ccacaacaag tttctagcta 60
tatgaactaa tatatgtgtc acttctaata tttctatggc tggtagctttt aggtaaagac 120
tggaagacc agcttgagc aaatatcaag agtttattgg aatcaggtaa gctaaaagct 180
aatagtcttg gccattctt tttcttatcc atgcacctt atgtacttga gaatccctaa 240
acatacatgn taacaataat tttccccata tgtaaaataa cttgacaccc tcgaacttct 300
canagtcatt ccaatttcta ttcgattcgc cattgttact ggtattntct acagattatg 360
ctcctgtcag tgttttgtac catcaattgt 390

<210> 30936

<211> 584

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30936

tcgcgcctca cacngccata taanactttc gttgtaatan ggtcaagtag acatgtactt 60
gttttaccac tacacaccna nccccacgaa gcacatttga agcccttgag ncctnngtga 120

aaaccagtca agaacacgag gaacctcgan agacgacctg caagcatagc agattgtata 180
 ggaagttata caggctgaca agagagaggg acacgaacgg aactttgaat agcatctcag 240
 agacattctt cacaaagtac aacaagcgct acacacgctt ctagttataa acaaggaaac 300
 acccctgaca atcctactga agaaaactct ctagagaagc attatttgaa tgccagagct 360
 ctacgacaca cacaccatct aaaactaagc tcacctccat gagaagctag agccaatcta 420
 cacacacccc gttaaataac taagactcac cttccttaag agagagcaag cgtagagcat 480
 taactaccac accctctatc atcagctaca gtcaccctc catgacagaa aaccatgata 540
 agtcactaaa aaaatcctac tactaagaac actcacaagc tccg 584

<210> 30937
 <211> 415
 <212> DNA
 <213> Glycine max

<400> 30937

catcactcga cccggatcct tagagtcacc ggggtcagc tttcttaacg gctgaatctc 60
 tccttaggct tatgttacta tttttatggt tcatttcttc ctgaggttgg ctgagccata 120
 gatctgaatg aagagacact agtgattcat gtatcctttg tctccttttc cttttctcca 180
 agacctacag cgtgtcaaga aggatataaa agagaaaata ttggcttcct ttggctatat 240
 gtattcctta gttagttggt cctaccttat catgtgatcc tagatcactg tgaaatgaat 300
 agaaggacac ctattgaatc atattttgaa tactagatca attctgctct gtctgtatat 360
 attgtggcat acgaaattat ctattgtgga atttccacaa agccttggtg ggatc 415

<210> 30938
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30938

acggacctat aaaactaagc ttagaccgag cactcaaatc taggatctaa aacccttcta 60
 tntagggatg ttcaagggtg agaaggaaaa tgaaaatggg gtaaatnngg agcaaactct 120
 cacctcaaaa aagtctatat catcaatcta aacttgctca aactggtttt accaagaana 180
 ctctaccgaa tcaaaagttg actcctcaac acccatattt taccctagaa atggctcttg 240

Figure 1. The ^{13}C NMR spectra of the polyimides 1a and 1b. The chemical shift of the carbonyl carbon is 165.0 ppm in 1a and 165.2 ppm in 1b. The chemical shift of the aromatic carbon is 120.0 ppm in 1a and 120.2 ppm in 1b. The chemical shift of the aliphatic carbon is 20.0 ppm in 1a and 20.2 ppm in 1b.

ctgggttttct gcaaaaaaac tctctgaatt acatgatcaa gagtatgtat atactctagc 300
 tggtagctct aggcacaaag gctatacttt ggtgccaaga ccaattcagt tatgacaact 360
 gaattca 367

<210> 30944
 <211> 389
 <212> DNA
 <213> Glycine max

<400> 30944

agcttcttag tttcagatga tgcagatggg tttgtagcta cctcatgcac tectctaattg 60
 actatggcat catttctggc gctaaactgc tgggagttgg aggccatctt ctcaattaaa 120
 tttctggctt cagtaggagt catgtctcca agggctccac cactggcagc atctatcata 180
 cttctctcca tattactgag tccttcataa aaatattgga gaagaagctg ttctgaaatc 240
 tgatgggtggg ggcaactggc acatagtttc ttaaactctc cccagtactc atacaggctc 300
 tctccactga gttgtctaata acctgagata tcttctctga tggctgtggt cctggaagca 360
 gggaaaattt tttctaagaa tactctctt 389

<210> 30945
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30945

ggaagctcaa ggaagagctn gaagaagttt tggctattac ttgcccact cctttgagtg 60
 acatttgat tggttgttat cttgattgat gcatcttagt acatttgata tctgctttgc 120
 atctgcac atcatgggta gcatcgagaa aagtttctaa gttaagaaaa tttcttcaga 180
 ggtaaaactc tctatttaata cgatacagag gtgtcggaat cgattcaaca agctgggtga 240
 agcttaaaga gttaagtctc atatcggttt aatccgatac aatagtactt taattgattt 300
 cactgctgtt agaccatgac tgatctttnt caggagtctc aactttaatc aattaccagt 360
 ggattaatcg attacttctc tctcgttcaa gtgttcaaag gtgaactata acactttaat 420
 cgattatat 429

<210> 30946
 <211> 400
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30946

agcttgtagt gaggtttgac tacaaaaatt cattggtttt tctaggattc aaaggtttag 60
 attctaagag agcacaagtc ctgactaat cccaatgac ttttcttggt ttgtacaaat 120
 agccttctca ctattgcctt ttcttaagtt gcttttgacc ttattgtaac aacataactt 180
 attttctctc tttttttaca ttcaacttat ttgatgtgtg tcttgatgct taactttttt 240
 cttttcattc ttttcaactt ttctcccca aatttagagt aaatatgcct tgaacaatat 300
 gctctcctag aatctaaaca aggtattagg agataatcat gtaaagttca gggttcaatt 360
 catgacaaat caataagctn tataacaacgc agcaaaagat 400

<210> 30947
 <211> 90
 <212> DNA
 <213> Glycine max

<400> 30947

agtacgtgaa ggaactcttg aaaaatttta gatggacgat gcaaatatat gaaactctat 60
 acatccacca ctatattaga ctagatatga 90

<210> 30948
 <211> 400
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30948

agcttggtgct gtatgggctg atcaatgttt tgacaaagta atagggatag aggttttaga 60
 tgaattgata atttggtgtg taattgaagg ctcttgagtt ctgtaactcc tgttggtgaga 120
 ggtaatggga gcacaggata tatgtatata aaggcaagtt gtgtgggcct acgggaagta 180
 tgatTTTTTg tcgggataga aattggaagg ggatacattt gtaggtttgg gttattcttg 240
 tttatgcac acttgacaaa acacctataa cctgacactt ttatgtgtgt acaagttctt 300
 aatggattcc anacatccat gatgatatga aaataagttg aattatttat ccagaagatt 360

gtgaaagatg tttccctgac aatattcttg ttattggttt

400

<210> 30949
<211> 429
<212> DNA
<213> Glycine max

<400> 30949

tcaagaataa tgggtctcatc aaactattta tttctcgaag ggaattctat aaataggcct 60
cctatttttta atggcatggg ttaccattat tggaaaaccc gcatgcaaatt ttgtatagag 120
gctatagatt tgaatatctg ggaagccaca gaaattaggt cctacattcc cactatgggt 180
gcaggaaata caccataga aaaacctagg gaagaatgga gtgaggagga aaagaaatta 240
gtttaatata atttaaaatt caaaaatata attacatatg ctttaggaat ggatgaatac 300
tttagggat caaattataa aaatgcaaaa gatatgtggg ataccctaca ggtaacacat 360
gaagatacaa catatgtaaa aagatctagg ataaatacat tgacacatga atatgaatta 420
tttagaatg 429

<210> 30950
<211> 403
<212> DNA
<213> Glycine max

<400> 30950

agcttaatcc cttgataatt gagggtagga gatttgcctt ggattcagct agggactact 60
ttccttagca cccttatgtt caatatgttg gataaataaa aatagttttt ttttgctata 120
tgcatgataa tttcgatgct agttatcaca caaatgtatt atacataagt acctatcaca 180
taaagagtgg ctatgcaatt tagaatgcat caagaagttt tagattacgt ggctacattc 240
tttggaaaca aaggcatcgc atggaaaaat tactacatac ccatatctaa tgggaatttc 300
tattttccta cttggctttt gtgagggaga tgtcaccaca cgttatgcag gatgggtggaa 360
gcagtcaata ttgtatcatc atcgtgattt tgcaaaaaat att 403

<210> 30951
<211> 412
<212> DNA
<213> Glycine max

<400> 30951

agttgggggc acttataatt taactcattg atttgtcaag aaaactattg gaaaaacgag 60
agtggtgaat gatttttggg ttttgtcttg tatgaacatc ctatagtaaa atttacattt 120
ctctctatat gattaacttt ggattttgag tcatacctta tccaaaacta ggcatacatg 180
actttttata ctggtaaagt ttataaaatt ttgttatatg ataagtataa gttatgcaat 240
agttttataa gtactctata gaaggaaaaa aatgtaaact tgagttttta tataaatttt 300
taatcagata aatatattta tgtataaatt tggtttgggt tggattagat tgaattttta 360
aatgaaatcc aaaatctgat tctatccaaa acatatgagt ttgttaaatt tt 412

<210> 30952

<211> 399

<212> DNA

<213> Glycine max

<400> 30952

agctttgaga aacaagtgat catccattgg caagtgatca tccattggca tcatcaaaac 60
attcagcttg atccattatc tacattatgt tgacaccaga gccatcgcca actaattact 120
aatcagtacc atgataggga ttgttacagc gtagattttt gcacaatagt ccatgatcca 180
gttattttgc aacggaataa ggtgtggtct tggcatgatg agtactaggt ccacttggac 240
attttggata gccatgaagt ttatagcaaa ccatggcagt gttccctagt ttctcatagt 300
attgacaaat aacattcttt gttatgcctg aaaatatgga tctgccatat gttgaagtgg 360
tttgttaaac atccgcaaga ccatgttttg atagtatgg 399

<210> 30953

<211> 153

<212> DNA

<213> Glycine max

<400> 30953

tatgttaatg ggtcttaaga ggaaagctca ctaacacaca ctcaaactta cttattaaac 60
atgctcatga aaactatttt ttctcaatta aaataaatcc cttttatttt cctacaccaa 120
taacccaaac tagaattaat taattaatta att 153

00424106-101500

<210> 30954
 <211> 396
 <212> DNA
 <213> Glycine max

<400> 30954

agctttttaa attttgaaat aaaatgtgaa aaactaatat attagaaata agatttgcta 60
 aaaaaagata aaaaaaaac aattagatct taaaaaatga acattaaaaa taggtacaga 120
 agaagaagaa gaagcatctt caagaaaaaa aatacatcag catcttcact cactcaccac 180
 gaaggttccg ttctctgtga atctatatat ttctagggca cgttacgtgg aacgaagctc 240
 ttcactgaat tcaatcgaag aagaagaaga agatggattg cgttgctggt tcgagtttgt 300
 ttccgttgca tcgttgcaaa accattcacc tggttggtaa tattcactct cccctcttca 360
 actcaccaca cacacaacac aattatctca ttcgca 396

<210> 30955
 <211> 372
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30955

tgcacacaag tttctccttg cctagcactt canaaccttc tggttgggtc atataaatgt 60
 ctctctctaa atcccatgc aagaatgcag ttntaacatc taactattcc aagtgaagat 120
 tctttgtagc tacaatgctt agaataactc tgatggtagt catctttaca attggagaga 180
 aaatctctat gaaatcaatt ccttggtttt gctgaaatcc tttcaccata agtctcgct 240
 tgtatcttct tctaccatca aattcttctt ttagcctata gaccactta ttctgtaaag 300
 ctttctttcc ttctagcaat ttaattaaag accacatctt attcttctga agggatgtca 360
 tctcatcttt ca 372

<210> 30956
 <211> 303
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30956

agcttttctg caactttctt ccattgatct gggttgatct taacaccgac taacctacct 60

0044106-101500

gaaattatca gccacatatg ccacggtggt gccctagtgt cttgagaatc atgacttate 120
 gcatgatcca gtatgccctt atagtagatc gagctcttag cggatggact aattgtcact 180
 aagatacttt aagtctatac tcacacctta cccaagagat gaaaaccttt agttaccatt 240
 caaaaggagg ccatctgcta caatgtntat atatacatca tgtacctgat gttctacagt 300
 tga 303

<210> 30957
 <211> 419
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30957

ccagttctgc gcaagctgcg agaattttac ggtggtcttc tgatagcacc ctgatgtact 60
 atggcagtgct ctctgtata gcttgccctac ctatccgcc cgctgagctc tgtagagaca 120
 ctgggtggtg agcagctttc tactatgagg cactactatc agttgcgaac ctatcgcaaa 180
 cgtggagagg aacatgactg aaatcgtgcc gtaatcacgt ggtcataaac catcactcga 240
 aaagaggagc tatgcgagg atcctctgtc tcatccgaaa ttgtccgacg gatgcgctgg 300
 gataaggaca ttgtcgactg ctngtcacac gagctatgta gtagcgaca taacagggtga 360
 gccgtgcatg gattggtcca caagatgctt tcgagaattt gactgcctga acgcagacg 419

<210> 30958
 <211> 310
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30958

aaagttcccg atcaaagatc ggaagaaagc aaaagaagaa aatttccgac caaagattgg 60
 aagaaagcca aaccgaaaag aaaagaaaat tcccgatcaa agatcggaag aaaatgaaag 120
 aaatatgcag aaaggtcttt ggaccaggca atatctgaac aatacagaat tgtaaccccc 180
 aaataaggaa ataaaggaaa ccacgaccgg aagtggctct ctccctttga tcgccaacca 240
 aaatcctgtg cgctagcgac tttctcacc cgactaaac anaaacagaa aaagaaaaga 300
 cccaaacact 310

<210> 30959
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30959

tctcaaggaa gttntcttaa gaaagcttct caaggaagct acctagtcta taaatagaag 60
 catgtgtaac acttggtgaa actttgatga aggagagtct tgtgagacat aactcaaagt 120
 tcaacttctc tccctttttc ttccttcaat ttcgtgctcc cccctctctc tttctctccc 180
 tctttctttt cctccattga agcctcctct ccaagcttct tatccaaggc tcactcttgg 240
 ggtgaatctc cttcttccat ggcttattcc ctagtggatg ggcctcctc tcacctcttc 300
 tcctttgtct tccgcttcat ctctatggtg gaaaaccacc attaaaggac ctcatgaag 360
 ctcanagatc cagcctccat agaagctcca caagcaagct tccatcaagt ggtatc 416

<210> 30960
 <211> 393
 <212> DNA
 <213> Glycine max

<400> 30960

agcttatgca tggattatgt aattatgaaa ttgagatgcc cgaagaaaca ccatttccta 60
 gctaaccatg cattaggtac catgttcaat tattttgttt ttgagtgaac cggttttatg 120
 atcccaacat gggtggctcg tgggtgcctaa cacatgaaac taagaatgta gtgtgaagtt 180
 tcacgcttcc cctttttttg tttttgttat gtagaggaaa acgcaaggat gagcacacat 240
 gaaaacaaat ggtatgcaat tttgcagatc aaaaagtttg ttgaacgcat atgcatgatg 300
 atgccatgac tcatgcaaaa tgtgaggctg gaatatgata acggacaaat gcaggatatg 360
 tccattatga tgttatgaag agatgcttat gcg 393

<210> 30961
 <211> 403
 <212> DNA
 <213> Glycine max

<400> 30961

tatgctacaa acatttataa tagaccctct cagtagcatt accaacaaca gcagaataat 60
 tatgatcttt caagcaacag atataatcta ggttggaaga atcatccaaa tctgagatgg 120
 gcaagtcctc cacaataaca acagcctatc cctcctttcc agaatgttgt tgggtccaagc 180
 aagccatattg ttctctctcc aatgcagcag cagacaacaa gcagctgagg ccccttctca 240
 accttccttg gaggagttag tgaggcaaatt gaccatctag aatatgcaat ttcagcaaga 300
 gacaagagcc tccattcaga gtctaacaaa tcagatgggg atgatggcta ctcagttgaa 360
 ccaagcttag tcccaaaatt ctgacaaatt tccttcacaa act 403

<210> 30962
 <211> 383
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 30962

agcttacatc acttggaag ataaagatat agattcatca ggagattcag aaaatgaagt 60
 cgtgaatctg agtctcatgg ccaaaaacta tgatagcgaa gaagaggtaa catcttctaa 120
 caataaccta tctatttctt ttgatgaact tcaagatgaa tttaatgact tgcataaaga 180
 atcagtcaaa cttgccaac tagtttcagt ttctaagaaa ataatttcaa atttagaaaa 240
 agaagttatg aaattaaata tagaattaga aaatcttaat actgaagtca aaacattaaa 300
 acaaattgat agaaatcaat cttctaccat accagatatt aataaagtat ctcactcatg 360
 tanatgttgt gataaattta aag 383

<210> 30963
 <211> 413
 <212> DNA
 <213> Glycine max

<400> 30963

tcctcggggc cattcctgcg aaggcaaaca tttggaaagt tagttttacc agtgggacgt 60
 tactctttaa agcaaaaatg gaatataacc tcctcccata aatacaacaa tcaatgtaaa 120
 tttagagtaa gcttatgctc atacttctt acaaatgttc tcttgacaaa gacattctat 180
 taaccgaaaa aatgcacca tatacaatca aggagctcc gttacctaga ttatttacac 240
 gtacttccaa ggtgtatttg ttacttacat cccacacatc tccttggtta aattcacata 300

agcttctact tatgtggcag ggcgggcttc cttcaccttc ttgtctccaa cgcgaacttt 60
gaccattgtt cttccttccc gcaatgcttc tcttcatgtc tgcttgagtg ggcttatagc 120
ctaaaccata cttcccaaga tttccttgag tatttatcag gctagtattg cgcgcgttgt 180
tttttctaa acccatcccg ggttcaaaac cgttcccaa cataactcgg gccatcatta 240
ccgctgcac ggacagacaa agttgcccaa agaggagtc cacggaggaa atgctgacca 300
cctcaaaaga ctggaaagca gtttctaacg attcttctgc ggcttcaca taaggcatgg 360
aggatgggca gcttaccaag atatcttct cgcctgacac 400

<210> 30967
<211> 296
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 30967

ntgaggggtgc gtagccacc atctnttcat agtagagtat cgataatgtg tctaccatca 60
cgatcatcgt ctccttttcc atcattgggg gtaccacctg tgccgtcaga tccctccacc 120
ttttgggcgt gttctttgaa agatccgtac cccttattgt aaatgttatg tagttgcac 180
ctatacgga ccatatccga attgtactga tactgactaa caaaggcaac cattatgtcc 240
atacaataat ggactcttga aaagtgaag ttagtgtacc atgtaacagc tacccc 296

<210> 30968
<211> 375
<212> DNA
<213> Glycine max

<400> 30968
tttctgaccg attgtctctc aatatgcacg ctccattcag ataccatgt catctcaggg 60
gggtagataa ggaattgggtg cccaatcgtg aattctgtaa agccaactac cccatcaaac 120
accttatgta tacctgtaag catttgatc tctattgcct cttcaacaac tgtatgatct 180
gtgagaatta tatatgacac caattctcta cccggtaac tcagtaatat ctttctactc 240
cctatatgta tctaccctta aatgacctaa attccatcgg ctatttatta aaaccgatcc 300
tttctggaag gttattctcc ctctttaaca caatcgcat tccctccccg catttaatac 360
cataatcctc gaccg 375

<210> 30969
 <211> 151
 <212> DNA
 <213> Glycine max

<400> 30969

agctttttatc tgataatata taagattcag ctagcctatg aataggcgta gaaataatcc 60
 taggcgagtt atattttgat agatctgatt ccgagagtca tatgtgtatg atgtgactta 120
 gagtcgtcta cctatcaata gcgtctctaa g 151

<210> 30970
 <211> 114
 <212> DNA
 <213> Glycine max

<400> 30970

tagcatgtac tatgatcttt gttggcgttc atgaagacca ttgtccgaaa gtagtttagca 60
 ttgaaaaacc tcgaaaccat agcattgggg tgagaaataa acctccaccc ttgt 114

<210> 30971
 <211> 320
 <212> DNA
 <213> Glycine max

<400> 30971

gctaagtgct gagttgagcg atagtgtttt atccttgttt gttcgggctc gttgaagcct 60
 tatctatcat tcccgacatg ttgacaacct gtcgagagct aatagagtat gctggacata 120
 aatatttgct tacaatgtcc aatgcaatcg cgccttgccc ataatgtggc gtatctataa 180
 tctaatatct gccaatacat aacatatttg aatacattgg aatattagtc caattataga 240
 ctatctgttt ggagggggagc ccggctacta acggtcacac tttcactttc ctataacaga 300
 ccagacccgc aagattgaca 320

<210> 30972
 <211> 565
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

005707 3072460

<400> 30972

ctgccacaca ccntggtaca tcnntataac atatatgatc atataataat tagcacaccn 60
cacaccctnn tnnncnncccc caccagnaca gcnnccttga aaccctgttg anacctagc 120
tannacngna cactatngaa tactgaagct taacaagntc atctatggat tgaaacaatc 180
ctcccgcta ttggtattac aatatccaga aggcatctcc tcattcagcg ttgaagagaa 240
tgtcccgat cactggttaa accacaaggg cagcgggaga aagaatcgtc tccttgatt 300
atacatacca tgatatctta ctgcgacta atgataaggg aatgctatat gaggcgaaac 360
aatctctctc aaagaactgt gataagaaat atatgggaga ggcaatttac gcataggcaa 420
aatactcata acaaagaact cgaagcattg tatggtgtgc cacagaacct atatcaacaa 480
ggttacagag aatacaaaga aagattgtca ccaagtgaac ctccaatgga aggtgacaac 540
ttcgttgaag catgccata atgaa 565

<210> 30973
<211> 294
<212> DNA
<213> Glycine max

<400> 30973

acttatcaca cggaagtccg attggagtgc ataatatatc gagaccctca atattgcaaa 60
aggtagtcct aatgaaagat aaatggggat aacttttttaa acggaagtct caattcaagt 120
gcatacaata ttcggaagct cgaaaatgaa caatggatgc ttctcgagaaa attaaatggt 180
cataacttat cacacggaag tccgatttag ggcgataata taccgagacg ctcgatattg 240
caactcgga gcaactcaaga aattcatgtg gtgataactt atcacacgga agtc 294

<210> 30974
<211> 563
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30974

accaccacc cactgcactc acctaatata gctgacatgt tgcacaatcg tacataataa 60
cacaaaaacn naaaaanaaaa aacgagnnac attgaagcca ttggaagcca tgtagaatcc 120
atggccaaca cnagctcnac acccgagaa ccacctgagg cgacctgcag gcatgctagc 180

atttgtgagc tttttaagtc tcagagaaac gagacaacgt gataccttac gacaggaact 240
 ccgaaatact aatgagaact aggagtatca cctatcccac aacgacgagg tggaaggcat 300
 ggccagcgaa gtaccaccat acacagtagg agaacagaaa ctagtcatgt actcaacagc 360
 tataatctat gaacaccata ttatcccgtat tacacggccc agtacgcggg aacggacaca 420
 caagcacaag caaacatcac gtgtgtcagc taacaacaac taaaacagtg ttagactaga 480
 taagcacatg agagagacat gaatggaaaa gaagaccaac gccataaacc ctgagtgtaa 540
 gaaattagac aaaatgacga acc 563

<210> 30975
 <211> 521
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30975

nggatgccct ttgtgatact tatttgaaac ccatgggtcaa gcncncnagn ggaactctat 60
 aatagttgac acttcacagc atgcaaagct ttcttgtcta tattgtataa antatttatt 120
 cttttgagc atcatctgac ctcaatggca tatactctta ttactgaata tcccacgtcc 180
 tactcggttaa tcaagctatt cgctcaacac ttaatgtcac tatcttctga tatctgaatt 240
 tctcactcac tgtaatttga tgatcacgca tctagtgcac cgaatttctc atcaacaata 300
 cgtgtcctgg ctgccactat tatgtagata agatgctcat gtcaggcttc tcgagttaat 360
 gtactatatg ctcccatctc ccccgagttc caaagatcaa ccatctcacc cccgcttact 420
 ttgttatctt tggacttgac attctcaact cactgcacga agtacacacg atcatcctat 480
 ttctagtttt tgcttaccta tgccctactt attgtaccac g 521

<210> 30976
 <211> 287
 <212> DNA
 <213> Glycine max
 <400> 30976

taatacatgc aattcatatg atgaaaacat ttatatcact aaaatctata gtaactaatt 60
 aatttaattc tatacattat tacatagacg gaacgtttat accatgcac aactaaataa 120

accttttttag gaaaaataaa aataaataaa agtgaagaaa aaaaaagaag agagcaacaa 180
aagacatccc gtgttcgagt ttttcattga taaaaactaa caacgtttac aagttattct 240
aatctcaaaa aaaaatgaat cagcatagag cttcacatca ttattca 287

<210> 30977
<211> 204
<212> DNA
<213> Glycine max

<400> 30977

ttttggggac ggcaacaaac ccggaatggg ttttaaggcca aacaactacg gaccactta 60
cctgggtcaat gcctaataa atcgagggaa gtatgggtta agctataatc ccactcccgc 120
cctatatgaa aagaagcatc tttggaatgt agaacggatg ccaaagctcc cagttgtgac 180
aagaatgtga agggatccca ccct 204

<210> 30978
<211> 406
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30978

agcttcatgt ttataggata aattattntc ttttggttatt cactgacctt aagccataat 60
ttaaaaaatga atattcacca aactgttaac aagcattgct caaacttaat gttatatttc 120
tgattctgat ttctcactag tgaatttgag gatcagcagt agtgatgaat tccacagaag 180
aagagtgtcg tggctgagag attatggaga aagatgctca tgtcagggtt ctcgagttaa 240
tgtactatat gctcccatct ccgtacgagt cccaagagat caaccatctc accctcgctt 300
actttgtcat ctctggactt gacattctca actcactcca caaagtacac acgatcatcc 360
tattcctagt ttttgctnta cctatgccat tactattggt accatg 406

<210> 30979
<211> 423
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30979

tgtgtcattt tcttaatcag aaaatggagg catgcatact tcaagtttct ttatgacttg 240
 aaagaccaac tttatgtcat ggtggccatc aaaatgaatt cttttcgttg aggtcggaca 300
 caaccatact ggtaagtaaa acctttgatt agttgaaatt tcngttttta ttatacaaga 360
 taaacataga ccatatttta acat 384

<210> 30982
 <211> 268
 <212> DNA
 <213> Glycine max

<400> 30982

agcttattca gattctgttt tacatattcg agaattacta ccgggcaacg tgaaagtctt 60
 aattcagtgg aaaaatctcc ctcccagtga acatagttgg gaatctgcgg ctaaatgaca 120
 ataggttatt ccgacttatc accttgagga ctatgtgagc cttttaggcg gaggtattga 180
 ttagaataag cataatccac acatcaccaa tgtgtacacc cgcataaatc ccatgggtgtg 240
 caaaccaccc aacatttaca acccaccc 268

<210> 30983
 <211> 426
 <212> DNA
 <213> Glycine max

<400> 30983

taggagtga ccattaaggc attgaccaac agtatgccat cttcttcaca gtgatattga 60
 gggggaaaga gagtgtgcgt taattaaatg ttttgaagtt ataaaccag ttacatgtc 120
 tctcattata tcggttgagc tatcttaaaa gaattgaaga tgttttaaac tacacaacga 180
 gacatctttt ttttcatttc tataaattat accaatcgcc attgtaatat caagtttatc 240
 aagtgatctt actaagattt tgttaaaaat aatcccacat cgagtaattg atgaacatga 300
 taagtgccta tatagctagg tagaccaccc ccttatgaac cggtttttaa ggtgacctt 360
 cggatgcctt tgctacacta taaaatctga tatggtatca gagccatatt caacagccct 420
 gactcg 426

<210> 30984
 <211> 400
 <212> DNA

0042105-10159

<213> Glycine max

<223> unsure at all n locations

<400> 30984

agcttgtaac tcataattnt tagttgaaat tgtcaatttt acacatgcaa tcttaattct 60
caacacactn tttggatgag tcttccaagg attgtgttgc cttctctaac tnttcttct 120
tttccagcga taaggtaaag ctacaaaatt gagtcttcca atgtttgata taagttttgc 180
aagaccatct ttaattcgaa taagtggctt aaagggtgaa atgcacagtc cttccaagcg 240
agcaactcan aggtgtaaca ccatcttaga atttcgtatg agcatcttca atgaaaatgg 300
aagacttgaa cgaaaatggg tggcttgctc ctcatgttgc tgggaataga taaggatcta 360
tataatgagc acaatgtatg aaggatggaa naacttcaat 400

<210> 30985

<211> 433

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 30985

tgtgcatcca ataccctgat gaggatgtcc catatgttct taaaactgga ctaattcatt 60
tgcttccaaa gtttcatggc cttgtagggtg aagaccgcga caaacatttg aaggaatttc 120
atattgtctg ctccaccatg aaacccccag atgtccaaga ggaccacata tttctgaagg 180
cttttctca ttcattagag ggagtggcaa aggactggct gtattacctt gctccaaggt 240
ccatcacgag ctgggatgac cttaagagag tattcttaga aaaaattttc cctgcttcca 300
ggaccacagc catcaggaag gatattctcag gtattagaca actcagtgga gagagcctgt 360
atgagtactg ngagagattt aagaaactat gtgccagttg ccctcaccat cagaattcag 420
aacagcttct tct 433

<210> 30986

<211> 397

<212> DNA

<213> Glycine max

<400> 30986

agcttgatt atagttaaga gtcacagat cactatacca ttaactatga aaaaaagtaa 60

agcacttaac agactaagca ctaggtcaac agaaaaacta ctacaaagaa atagagtcga 300
 acatgattag gatcaggatc atgcaccaat tcgacccaag aatcaaagaa taggcctaaa 360
 atgacaataa tcgccgaaca aagtgaacta caattcactt aacgcggaaa taaaaggctg 420
 caacatactt gcaacaattt gcgacgatta ctatgagtga gaaagactct aacaggatag 480
 ggagtatgtg atgcacactt atatatattg cgcactttca gaagatgaat catgggtggg 540
 agaacttaaa aggaagggtca gcaccccg 568

<210> 30989
 <211> 384
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30989

ntgagtccaa cattcagttt tatacctgtc gtcatacttt attttncgcc ggcgaccttt 60
 gcttgatgac atgcgacctt tctttggtcc ttgtgaggtg cttggcacc atcattaggg 120
 aatttgtgaa attccaggac atgccagaaa accaaaaaat attgatgcac aatccgtaag 180
 tttccgtgac acaccggaaa tcaaattggaa gcatcggttc ataattaagt gagattccgt 240
 aacattccgt aagtcaaaaa ggggatgatt atgtaattcg caaggttccg taacattacg 300
 gaaagaaaac aagtatcggt acgagaatcg taagtttccg taactttacg aacaaagact 360
 caccaaaaaa ggaaggggggt gaac 384

<210> 30990
 <211> 232
 <212> DNA
 <213> Glycine max
 <400> 30990

agctgtttgt cgtctcattc aattgctgat tttccagcat gctctgaata gacttgcgca 60
 tcggttttaca catcgactac gcatcgatac catacctcta tacactggcg gatgacgcga 120
 ggtaattacc cttctacgct cattgatttc agaaaagact tcacgatcta gacgaacctt 180
 ggcatgccc acttttcttc tttgtcaacc actgccatct tactccaccg tt 232

<210> 30991
 <211> 392

0943406-10590

<212> DNA
<213> Glycine max

<400> 30991

taagcttgta taaattcttg tgatgaagca catttttgaa tccatctcta ttatcaaata 60
aataaaggct gaagcacagt gcttcatgat atttcaaat catggagaat caaactttcc 120
atgatgacca tttgaacaac tatatttctt tggataaaac tttttgcact aacagattag 180
tcacttaaaa gtatgatgga ccaattggca gcagatccag tcaaatgtgg aaatacttat 240
ccaatacctg ccataacttt aaaattatat acagatcatc actggtgttc tcccaaacc 300
gtgcatttcc aattgcaaaa caaccacatg gcgatctaaa aggatttgag tccagaggag 360
ctgcatcatt atacacaatc acatacatca tg 392

<210> 30992
<211> 563
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30992

ccctccgtca tctcgacat ctnatacat anagaaagac tagangtata nancgctcta 60
tagatgaagt acacttaaga aancnnnaaa ccagaggcaa cggtgaacc tttgangccc 120
atgtgcance ccaggcgata ccagctcgac acccgagat cctctacacg catccgcacg 180
cttgcaant agtatgaaca tggatataggc catctagaac gtgccaacgc atgccatata 240
cacgctgttt cgcttacgaa tcaatagcca gaagaggata aagcaccgaa atgaacaatc 300
tgaaacataa agtcactgaa ccaaaataga tacctacaca aatgggacaa cgcaaagcta 360
tcaactgccag actgagaagc aatgtttgat aggaggctac atacatacgt tttgctctta 420
ccactcaaac tgaactaaat caccaatctt ttctatgact cacgcccac tacataatca 480
aaaacttaaa cgcacacac ctgcctccgc atgactcaac gtcatagct aaacaaagaa 540
ctatcatcgt ccaattaata acc 563

<210> 30993
<211> 83
<212> DNA
<213> Glycine max

<400> 30993

gtggaactgt tcctcaaggg attaaaggaa gagattatca ctaacgtgag gtttcatgaa 60
ccatagaact agatggaagc tat 83

<210> 30994

<211> 368

<212> DNA

<213> Glycine max

<400> 30994

agctttcatc tagcctatat tatacaaaag tggtacaaca gaacctaacg gtatctaatt 60
atatggggcca tcaaattctat catgtgttga cagtaattga ttagcccatg aatttcctcg 120
ggggctgtac acacttcaac gatggctttt gctttggcta atagtcgcgg gaggtcttga 180
cttccattca aggtcaaggc gaacctatcc atccacatag tcgcttcttg atgcaatgca 240
tcaatcacct cctctctgct tctttttcgg tgtacacttg tgcaaatcc tctattagct 300
tttgttcatg ggtcatagac tgggtcaact ctcccttgta ctgcctatg atagctagca 360
tgctttgc 368

<210> 30995

<211> 387

<212> DNA

<213> Glycine max

<400> 30995

tcaaagatga ggtcaaaggc tacattctgt gtcaaatacc tgtgtcttaa cattaagggc 60
tgatgggtat ttcgggttct ataaaaaaga cacatatttt tgagattccg atcacgcaa 120
tgtgaccggg gttcggtgaa tgccgtaaaa acaatctcaa tgttataaaa agataactct 180
taaaatgtct cattctctat ggttattcaa aggaagtgtg tgatcacccg attacagtac 240
cctgcacata gaatacacta tgaggagctc aaactagtta cgagaatgct tagaactcaa 300
ggctacctca gggaaacttt gaaatggagg attctgagga ttatctccat ggaatcttct 360
aggaggattc tgaggatttc actctga 387

<210> 30996

<211> 383

<212> DNA

<213> Glycine max

<400> 30996

agcttgattg caagttgctt tgtctatatg catcttaatt cttctagatc ccatacctacc 60
attacaccaa gtgagactag atcccccttga acagaggtgg gtgagatcat tctagacaga 120
ccaattggaa aaatcttcac aagattgctt gaggggaagg tagccaccaa ttctttcatg 180
tgccccgtgaa accgcattga aatcaccaat gtaaaccctaa ggtccaagga agttgatcag 240
catagaagag agtcatgcc ataagattgc tcttttaatg ttggaggtgg aaccataaat 300
agtagctaca taacatgaaa tgttggttaat agaaactaca aaagacatgc tttgatcaga 360
gatagctaac atagacaagg aag 383

<210> 30997

<211> 311

<212> DNA

<213> Glycine max

<400> 30997

tgcacgtctt cactctctc aagcatttta tctcttctc cgctcagact ctttagcttt 60
gggagccaag ttatcccttg cgtactcgac ttcaaccatt tgagatagct gcctatgaca 120
ccttggttac ttccactaag ttctttatct tttctttctg ctttattcca ttccttatag 180
atcctctgga gtgtctttac attagcttca ttgaaacctc gcgtgatgaa aggcgcgatg 240
gtctcctccg atgggtgcacc tctcataggg taacctaaact ggcttatggc caacatggga 300
ttataattaa t 311

<210> 30998

<211> 377

<212> DNA

<213> Glycine max

<400> 30998

agcttgtaac tctgattcaa tgactgttaa aaacggtaa gatatactgc aaaattggcg 60
aactttatcg ctatctcaag atttcaaac atacaatgac tgtactttga aaaaaaatgc 120
ctaacaacta tctttagctt aaaattgcgt cagtagcata agaataatgc ttgtattcct 180
ttgttcacaa tgtaaaagat aactgtatac cacaacaaat atttcttagg cgaaaagaaa 240

gtgagcaatt cagatatttg aaaatgattt atgttttgat ttcctttcaa taaattacag 300
atgtataaac atatgctatc gcatagtcgc atgattcttt cttacgtaa gaagagatta 360
atattatggt ctatttg 377

<210>	30999
<211>	465
<212>	DNA
<213>	Glycine max

atganacctt	gtgattgaca	cnnttgcatt	acngnacact	tagaatactc	agcttcagaa	60
taattgggaa	accgtaattt	gtttaattca	ttcttattcc	agacttgtat	cattcgaaat	120
atattatcta	cctatatattt	attatattat	gccatatattt	aatgatggca	gaccattgggt	180
atcacagaaa	atgaattgca	cttttgcatt	acgggtgggc	ncattctagt	gatgagattt	240
acttacttct	acgtttgtac	cactacatgc	gggccttaac	agtaaagatt	attgtaattt	300
caaacactta	atgcatcttc	tatggcatat	ttggcaataa	ctatctctaa	accgaaagag	360
aatatatgtt	ggtgggcgct	ttcaacaccg	gagtaaattt	gtggtattaa	taagtacatc	420
atattaaaga	ggatctcata	cagcatcggt	ttaaaaaaca	gggcg		465

```
<223>      unsure at all n locations
<400>      31000
```

ttacattgtc atccattnga gttatcccag aatgtacatt tttgaacaca agtgaaatct	60
cccctttcaa agatgaaaga aacgaatggg caagtatcat caacccctcc ccaccacct	120
gacctcaaaa taacggctaa tggtaaacac tcctcttttg catttcatat tatacaataa	180
attctatatt tacaggagtg tattcatatt agagatgaaa aacgtacgat tacacaccat	240
gttaagagat gaaaactgaa cagactatga tactcactcg cgcatacatt gcttattact	300
tata	304

<211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31001

gggattcggc tggctacat attcagtatt ctatngagag acttgatatac tatgatggga 60
 agctgcactt gaaaatacca tcacattttc taaactggga tatttctgga tatacttcac 120
 gtggaatcct caagttgact atcaagtaat tctgtcatca gtgaggtgat ctactttggc 180
 attttctcat catttgaaag actcgcgatg caagctatat ttgaaagtct caaggatgta 240
 tcagcctcta attgcatctc attgtcctca gtagcaatc tatcgctttt tcacctaaaa 300
 tgaattaaaa actatctcgg acatgaaaca actcctacag caaagttgct atcaatattg 360
 tctagctcta tctatgcatt gagggactga ccacattgaa agtattgcgc ttt 413

<210> 31002
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31002

agctttgact tgagtcacat agagattata aatatatgac catggcatga atttcataac 60
 aatctttttc aacatctctt tcaataatca agaattctatc tttcaatctt ctctctcaac 120
 atcattcaac tttttctaca gaagtttggtg attcttcttc tcttcatctt tctaaaagtt 180
 tttgttcaaa actttttctt tcaagaaaag ttctttgatc aaaaacttgt gttattcatc 240
 tttttttatt cttttctcct ttgccaataa gaacgaagga ctaaccgcct gaattctttt 300
 gtgtctctct tctccctttc caagagaatt caaaggaccc cgctgagaa ttcttttgat 360
 tcttcccttt cccttanaca aaagatctca naggactaac cgc 403

<210> 31003
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31003

tgtaacgtgg tcatgttaac aagaaaatca tgggttttca tagattcaaa ctcttaggtt 60

ctacgagagc attcaccat tgcattgcta cttaaagaac cactttttct ttgacctccc 120
aacctttatt gacatgccac aaataacaga acatagaggt tctttttttt tggatgcat 180
ttgctttcag ctcatatttg cttttttttt tacgatgata ggtattacaa aagaatgtaa 240
atctgattct ctatgtatct gttactcata ttcttgaca taatttaacc aaaacactcc 300
cccaaatttg gaacaaattt gacttgatcc ataataatgc tctctatag cctaagatac 360
ggtgcacata gatagcattt acatttagct tanggttcaa tgacacatat cgtcacg 417

<210> 31004
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31004

agctttgagc aaattgtaat gacaataact ntatacacgg atgtccgggt gagtcccgta 60
agatatcgag acgctcaaaa tttagatccg aagctctgag aaaattgaat tgacaataac 120
tntatacacg gatgtccggt tgagtcctgt aatatatcga gacgctgcaa attgaaaacg 180
gaagctcgta ggacattcaa acgacaataa cttntactc ggatgttcga ttgaatcggg 240
taatatatcg agacgatcaa aattgagact agaagctctg agcaaattga gatgacaata 300
actttataca ctgatgtgcg gctgagtcgc gtgatatac gagacgctca aaatttagat 360
ccgaagctct gagagaattg aattgacaat aac 393

<210> 31005
<211> 206
<212> DNA
<213> Glycine max

<400> 31005

ctcgatatat taccagactc atgcggactt tcgtatataa acttattggc aattaaattt 60
tctcagagct ttggagcaaa attgtgagcg tctcgatata tgactggact cattcacaca 120
tccgatgaaa agattatttg cgtgagaata tgagacgagc ttccgttgct aatatggacc 180
atctctcgct atattgcgat aggcta 206

<210> 31006

<211> 391
 <212> DNA
 <213> Glycine max

<400> 31006

```
agcttcttat ccaatgctca tcttggtggt gaagctcctt cttccatggc ttattcccta 60
gtggatggca cctcctctca cctcttcttc tttgtcttcc gctgcatctc catggtggaa 120
aaccaccatt aaaggacctc attgaagctc aaagatccag cctccataga agccccacaa 180
gcaagtttcc atcagggttat ataagaggtc acatggtggt gtcctagaga attttctcaa 240
aaacaaaaat gtcgacatag ttagtttcga gagatctgca aacaccatgt attcaaatat 300
catgggggtg tggcatgtcg cttagaattt gatgtacatt aaaaatgtgg ctctcttctt 360
ttctcaaaat ggtgtttcat agttataaat c 391
```

<210> 31007
 <211> 414
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31007

```
tgttcggctt aattgtagtt tcgaatcgtg gagtagttgt tatcattatg ttttatgcat 60
cttttagtagt tggggtttga gacgggctcg aagatggaat gttatttata acaactttta 120
ctaagttatt cacaacctct ttaagatctt caagagttga tttcatattt gagatttctt 180
gacataacctg tgttatagat atgttttgtg tctcatgcat aacttttccc tctccaacag 240
ttgaacacta ctgcaaaaat aacatactac gacagttctt gagtacattt aaagaccatt 300
ttgaatcatc tttgaaacca acatcgttga aagtcttgac tnttgacgac ggttntcaaa 360
anatcgtctt agaaaaaagt atcattntaa gacggttctt gattaagaac tatc 414
```

<210> 31008
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31008

```
agctttctcc atttcatctt cattgtcaaa tcctccaaat cttaatccat acagttgacc 60
```

actcttgctc ctctggaag atatgaacac atctataacc ctccctattt acggaagatc 120
 cccacaagt tactaacatt catattcttt gggaatttag tgaaaaagaa tgtagtttca 180
 tcacctcccc tcccttgatc acctttccgt gctctacccc tccactctcc atcctctctc 240
 attttcctaa tgcacttagg gacacataac cccttatcaa gtaaaacaaa ttttaaaaat 300
 attcttggtt tatttagctt cttattctat taggattaat taaatatnta aaattcaata 360
 atattctaca tatttagcta aagggaacta tttt 394

<210> 31009
 <211> 424
 <212> DNA
 <213> Glycine max

<400> 31009

tgaactatca ggaaggatgg tgggtctaatt tgtagaactt ttataattcg acatctagta 60
 cgaaccctat ggtccgatga agacacaact catggtagac ttcctggaag aattcggttg 120
 gaatgaccaa accaccccag actggtggag cttctacgtt gacggtgcat ccaacgtgaa 180
 ggggagtagg gcatgaatca tctttgaagg ccctggaaat gtcactctaa agcaagccct 240
 taaatttaac ttcaaagcct caaacaatca ggccgagtac gaggcactca ttgcaggtct 300
 aaaactagca acaaaagttg gggccataaa gctctgatgc tacacggact cgcactctgt 360
 ccaggggcag gttgccaaact gataccagac caaagagaca atgttgctca agtactacca 420
 catt 424

<210> 31010
 <211> 206
 <212> DNA
 <213> Glycine max

<400> 31010

tagctgtttg aatctatata tggtttaaag ccagcttctc agcatgggga ccttaaggtc 60
 catggtataa tctgtggata atgggtatgat gataacctta tggatcaatg cctataccgc 120
 aacgttaagg ggagaaacag aggcgtcctt ggagtgtacg tacatgatat tatagttgca 180
 ggctatgagc ccggtttgct acatga 206

<210> 31011

agcattcaat cgcaatgcct tatgcatgcg atatctaaca agatgtgccc aatcaattt 419

<210> 31014
<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31014

agcttcccat ctctagcatc cctcttgaac ttcaaatacat actgatggaa cttccatatg 60
tactttcaatt tcctcagatt cctatagaac agagttgtgg gtatgttctg aaagtaaatac 120
ccaaagtcta ggccattctc atgcagagaa tcaaagatgg ttttttgagg ataccctttt 180
gctaactgcc tcttgatatg acttggtgaa ccatgagagg ttgctgagta cacaaaaagc 240
ctattggggtt gtgttggacc aggaattgaa gaanaccacc tgtcaaaaac agcaaattcc 300
ttaaccaaag cagcataaat cggcacagag tccggtttaa accctttcat gacagtctca 360
gagaggttgg gagacataga caatgc 386

<210> 31015
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31015

tgaactatta ggtaaaatcc tatccaataa tggctcttatt tacagaaagt gggtagggga 60
acttccgcaa aaaaatatag ctggtgtaat gggggttgat ttggcgtgta aggattgatc 120
agtgaagcac ttcaattgaa aggccttgac agttgtactt ataggtctga ttcccactgc 180
caacaatccc acttggtagg ttttattttt atggaaatat cactataagc tctcattgag 240
gattatgaca atgccctgcc ttggaatgga atagcctttc ctccccttaa aacttttgtt 300
tccaaaagaa gctgtcgtag attttggact tgtcacttat ggtaaagata tttataggta 360
tacatgtgct taanaggagt ntgacttngn attgatgaat gttgtanggg ttgaaattgc 420
a 421

<210> 31016
<211> 380

<212> DNA
 <213> Glycine max

 <400> 31016

 ttttagtgga tgtgatcatt aatagtcgct ttgaggtttg ctaaaataca accgattcat 60
 tctattatga ataatgaatt taaattttac ataagaataa ttatgatagt taaaaacact 120
 tataaagaga tgattaaaaa aatgtacaaa tctagaagat aatattaata ccatacaaga 180
 ataatagaat ataatatata tatatatata tatatatata tatatatata tatatatata 240
 tatatatata tatatatata ctgtgtatat atactcaacc tcatgcatat accttcattc 300
 aaattaaata ataacctata acattcgagc tgcgaaatct gctgctctca tattggatta 360
 tgaattcttt atctaaacag 380

<210> 31017
 <211> 352
 <212> DNA
 <213> Glycine max

 <400> 31017

 aatagtgaga aatataagaa ctattttaac ttctaattta ctctgattat tgatttatta 60
 cagtatgggt aaactatgat taactataaa gtatgatgga acaaatatga aaaggcttac 120
 cgactactgc tgattatgaa ataaatacac aacatacata gagtgaaatg agcatatatt 180
 tcatatgatt acaatgacaa ataagacctt attcatcagt catcaccaaa tgagcataat 240
 cacataaata aatggtcgga aatttgtctt aaattgaaaa ggaaaactgt gaggggccac 300
 gatcactgtc agctgtgggg aatgaataaa ttcaccatt cattcattga tg 352

<210> 31018
 <211> 167
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31018

 ttctccttac gcatctgtgc ggtatttcac accgcatatg gtgcactctc agtacaatct 60
 gctctgatgc cgcatagtta agccagcccc gacaccgcc aacaccgct gacgcgaacc 120
 ccttgcggnc gcatcgaata taacattcga taatgtatgc tataccn 167

<210> 31019
 <211> 291
 <212> DNA
 <213> Glycine max

<400> 31019

agcttttgat ataacttcta atgtgaatgt gatatgcgta tggcctaata taagctattt 60
 actctaatac accttacatg taattgtaca catattattt gcttatattt gatgtttggt 120
 gtttcttttaa atattggtga tccttagtga gcttgaaaca ttaacgtgcg gagtaaaaat 180
 tgcatttttg tttaatgttt caacaaaacc tttttttttt catttttttg gggggggggg 240
 gggggtgaac aaacaactga tgaaaatctc ctgtgataaa ctacaaaatc c 291

<210> 31020
 <211> 421
 <212> DNA
 <213> Glycine max

<400> 31020

agctttggca tctgagaggg gtcttcatgc tggtattgat ggtttctggg tggaaaatcc 60
 taatttggtt aagcctgaaa ttctgcagca ttgcaaagc agattcaaata taattgaagt 120
 tatgtacaag cactggagct tatttaaggc agaaattcta cagcatctgc agtctgtggg 180
 tggaaaaagg gtgggagtg aaatttaaata ggagaagaca cttgtttgac agagaccttg 240
 agatggcaga ttgtttccgt aatgatgttg ctggcaactg tatttacatt cacaaaaagg 300
 atgagtggat ctggaaaata gaccctactg gacaatatcc ggtaattaaa ggagagacta 360
 caaacaaca caaatttgat gaggataacg gtggcaatta atgacacatt atgcccattc 420
 t 421

<210> 31021
 <211> 393
 <212> DNA
 <213> Glycine max

<400> 31021

tgtaaccgag gcatgctaga cggctcttat attacaaga aatccttcca ttgtaagcac 60
 accacatatg aatgagttta tttttggaaa tatcatcatt aatgataaaa ataaagatct 120

<210> 31024
 <211> 401
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31024

 agcttgtagc tttcatgact tcagaaaagc caaagatttc ccccatccta gttctaaatt 60
 tcaaagtcta tcaaagctta atttaaaata tgtgtgcaact ttggaccaca atgtgtgcaa 120
 tttctgaaat ggggttcaat ttattataac aaatagtgat gtacataatg tttgtacaca 180
 tttatatatg tgtggcataa tgaattatac tcgtaaatga aatatactgg tttaggattt 240
 gttttatttc tgcatacctt agcaaaggaa acaactcttg taaatttgtc aagttaatat 300
 ccaaaagctt tagctacttg agacgcacga tctgcatatg ccaaaaaata ttatcgtttt 360
 ctccatcatc caaactnttc agttttcctc ctggttatatt t 401

<210> 31025
 <211> 423
 <212> DNA
 <213> Glycine max

 <400> 31025

 ctcatcactt tcagcaatac attctccac tcaaatagtc tccgatgccca ttcattatta 60
 tagccaccat tctgaccac ccgagaatca acttcaactga caaggatatac tttatccatc 120
 acaacgagaa caatggccag atttatcct tcatggtaat ttcacccttc catatatcta 180
 tccaaaatgc atatatgcac cattccccac ctttctaate atattaactg aaaaccaatt 240
 caccggacat aatgaattat gattttcccc catcatatcc ttccaccata taaaagtttg 300
 actatgagtc aaactacctt ccacatccaa tcaatagtca tacctgaaat gtataaaatc 360
 aaactatata gtgtgcttat ctatagagat tctcctcctc cacttagcta gaaggcttgc 420
 att 423

<210> 31026
 <211> 401
 <212> DNA
 <213> Glycine max

 <400> 31026

agcttatcat atgtagctga gatgattgga gttccaacaa ctagattatg aagaagtgag 60
gtctgcatgt ttgatgatta acaaaaaaga tttttgtttt atcaaaggga ctagaagcgg 120
gtgaacagaa acaggtggag ctactagatt gctagggaag ttatgctaac ttgggttgat 180
gagttccttg tattccaaag aggtatttaa agggatggaa actaggaagt gaatctttga 240
cactttcttt agtggggcaa ctcaaccttg gatcatcaac cttgattgat aaaaatgtac 300
tgatatgaca accgtgacaa ttattcaaga taatattttc acatcatgga ctgttttgct 360
tgctttaaga aatatgggtc ggtaagggtt agctttttga g 401

<210> 31027
<211> 430
<212> DNA
<213> Glycine max

<400> 31027
tacctgagtt aattgcctgc tttcatgatt tccacgtaat agggtaatat tagctgggta 60
tctgtgaaat tacagaaaaa taaaaggata tgtaagtttt caaaagaaaa aaaagaaaag 120
caccactgca aatgggtgta aactttccaa agtcaaatat gctagcaatg aaagactggt 180
tggaagtatc ataagcattt attgtaagat tgttgacttt tataatactt acttttaaag 240
ttatatcaag cacaatttct aatgggctga cagtgtaaaa atcttcacac tgaccgttta 300
gcaaacttat acactttgat accatgggat acaaaccaaa ttaatcttaa aactcttctt 360
acatgcctcc aaagctagga caaatttcaa gtgataatct tattaacaat acatgtaaag 420
ccaacaagtc 430

<210> 31028
<211> 329
<212> DNA
<213> Glycine max

<400> 31028
aaatcagcca ttattgaacc attatgaact ctccaaccac cgggaccatc tccggtggaa 60
gaatcattcc aaacaacagc aacaacaatc ttattttcaa aatgctattg gcccaagcag 120
aacatacatt tcttcaccaa tccagcaaca acaacagcaa ccgccccaga aacagcaaac 180
aggtgaagct ccttcgcaac cttcccttga agaacttggt aggcaaata ctatgccaaa 240

catgcagttt caacaagaga ccagagcctt cattcaaagc ttaactaatc agatgggaca 300
catggctaca cagttaaatc aacaacagt 329

<210> 31029
<211> 315
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31029

ccttgtccga aaagtcactt anaaccatth taagggtccaa cgccttanaa cggtcctctt 60
tgctttttatc gattaacatg gaccgttcaa aagcataaga tcaacacata actttaccgc 120
ttttgcaaga actatgtagg tctgagttcc tcatcacana tcgaggatac gtangagcaa 180
aagccccgct tttgtcgacc accccaagag atcggttaatg gtccaacgcc ttaacgtttc 240
tctcctttca aaaaccaaga gatcggttaat ggtccaacgc cttaacgttt ctcttctttc 300
aaaatcaaaa gatca 315

<210> 31030
<211> 392
<212> DNA
<213> Glycine max

<400> 31030

tagcttgtga agatgcttca atggaggaaa agaaagaggg agagaaagag agagggggga 60
gcacgaaatt gaaggaagaa aaagggagag aagttgaact ttgagttgtg tctcacaaga 120
ctctcattca tcaaagttac cacaagtgtt acacatgctt ctatttatag actaggtagc 180
ttccttgaga agctttcttg agaaaccttc cttgagaagc ttctttgaga aaacttcctt 240
gagaagctag agcttatcta cacacacccc tctcataact aagctcacct ccttgagaag 300
cttctttaag aagattccta aagaagctag agcttagcta cacatacctc tcttatagct 360
aagctcacct ccttgagatg agaagctaga gc 392

<210> 31031
<211> 413
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31031

tcagaccaag gcaactcana atctaggtat ctaaaacccc tcaatttagt ggatttcaac 60
gtttaagaag tgaaaatgag aatggggtaa atttggagca aactctcacc tcacacaagt 120
ctataacatt aatctaaact tgctcaaact ggttctacac ctaaaattcc accgaatcaa 180
aatttgactc ctcaacaccc aattttaccc tagaaatgac ccttggtttc actttgggtca 240
ctcatactcc tcatttgcac agtctaagct ttctcttaag tcctaaatga catttcaaac 300
taagattaac tcactttaac cccaattac cactgaatcc agatttagcc ttccaactct 360
caaagcctca ctctttttcc actcataaca ccacattctc actttctaac cct 413

<210> 31032
<211> 285
<212> DNA
<213> Glycine max

<400> 31032
cacttcttat gctcaaagaa gaatcacctc gatcagaaag aactacgcag gtctgatttt 60
ctcatcccaa ttgaggaata cgtatgagca tagggaaaca cccttgtcga cctcgctaag 120
agaaactata tacaacgggt ataaaggata taaatacata caacgggaac ataaaaaatc 180
aaagtcacgt ttgcacattc gattaaagggt tgccgtccct tgcgacggac gtgtgggggtg 240
ctaatacctt ctccgtgcgt aaatacaact cccgaacctt tcact 285

<210> 31033
<211> 476
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31033

cggcgccatg accccnattg agtccttgca ttacgtgaca cttaaatact aagcttgggg 60
agtctcgcg cccaccacat gatggtggtt ggtgtctgtc ggagcatcgg gtcctggag 120
gaatcctcct gacgggcaca gcgcggactg gctgctatct gcagccgcct atctactaat 180
gagccacccc tgcttttact tggcgattct ttttgggtct atgaacacgc aactcaccaa 240
tttctacca gacttgcgaa ctttccataa tgtcacgta ccttgcggaac taactaatc 300

atcccatatt gacttacaga gggttacgaa accgtcctaa ctgcgcaccg aagcacacat 360
 ttgattaccc gtggacccca gtaccatacc gattgtgcag caagataacc gtttgatcta 420
 ctgcacgtac cggaagctca catatagtct tatgaccggc ggcaagaacc tcgcan 476

<210> 31034
 <211> 321
 <212> DNA
 <213> Glycine max

<400> 31034
 agcttgtacg ccaaatcgtg actggccata tcccttgacc gatggagaga ttgcacttcc 60
 ttgccctctg atgcattacc ctagtcgagg ggaatttggc gagcgaagac aaatactcag 120
 aagctttgca atgcatgcta cacaagagac gatggaaccc tgggtcatct aggattgtct 180
 gagagcattc aaaggatctt attcgagact acccgatcaa gtactgtgag aggggggaaat 240
 gaatagagga gacaatttcc attaccctag tgtagttgta tacaggcaga cgacgaatga 300
 cctacagctc ctgatcacga t 321

<210> 31035
 <211> 407
 <212> DNA
 <213> Glycine max

<400> 31035
 gtgagaatgt gtctgaagaa gcacacgatg ttagacgacc tttacatctc tggaatccca 60
 atgcggactc tttcagccac tttccactat tgcatagttc ctacctccaa tatcaacatc 120
 atgactgctt tatcttatgc ttacagcaaa ttgcatctca ccgagaccat tctccaaccg 180
 agaataccca tgtgtatgag attcctccag ccacaatcca tccagagatc tggccctcca 240
 cactgtgcat taagagtgac attatgaact gattgccgac ttccaacggg atgtgtccaa 300
 cgagcctacc tgatgggtga ttacataacc tatgaattaa atatgtctag gctgactacc 360
 tatttggtc cctgtatgac attgtgatgg tgtagttgct aacaaat 407

<210> 31036
 <211> 269
 <212> DNA
 <213> Glycine max

<400> 31036

gcgctctgga caacaccgaa aacacccgga tatgcagtgg tcaaacgaaa aacaagaaaa 60
gcaccactgc agacgggtgc aaactttaca cattcaaagtg tgctaccaat gcaagaccga 120
ttggacgaat cataaacatt tattgaccga tggccgactc ttataacaca cactttttaa 180
gtcatcataa gcgccatccc gaacggggccg acacagtaaa aggctccaca ctgaccgact 240
agcagactaa tacgctaagga taccatggg 269

<210> 31037

<211> 377

<212> DNA

<213> Glycine max

<400> 31037

agctttttga actaggatgt gttagatcac ccaataacgc ggccacatac acagcttcta 60
gctattcgta ggacatttca agggccgtat ccacatgtca atgttacacg gtgatgtttt 120
taccataaaa tcaaacctca ttgctaacta ttataaatat ggcgagagta acttaaaaca 180
tttattgttg cctcatctta ataccaatct actctcgatt ttgctatata cgtttggttg 240
atgattgttt cctagagatt ttgatgtcta ttcttataga ttttaaattc ctcatatcat 300
attgagaatt ggccttggca tcgtgtgatg tgcccatgc agcagcaact tctgttttac 360
tcatacttct tactata 377

<210> 31038

<211> 601

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31038

cctcccgcat ctccgnctcg ctccccnct aanntntccc tctctctct atactctcg 60
tccttctctc gcannctna nntaaannnn nntanannca gcnaccnagc ggacantttg 120
gaaacccttg gtagatttgc agtacgacta gccanancng ngacactata nnaaactcaa 180
gcttggccac tttcatccag aactggtagg ctcanatctt cttctgtcta cctcgacgac 240
gagaaccaga tcctctctgt catcggagac cacacaagca tgcaagtaaa aggagaatat 300
attctaacia tagagcacgt acatcgatgc acaactctac actatcacan attatgatag 360

gacgataatg ccggaagagt ctctgcaatg agttatcctt gcaaacgcat acgtacacaa 420
gaattccaca aagttgacac cttaggtgta taacactcaa cactgagtac aagaggacct 480
actcgttacc atgtggggcc tcctatgtta tggaaactcg gtgagcacca cccagagggc 540
gtgccataca cttacaggta accttaccga gcctgcccgg aatgtctgtc ctaggaacgc 600
n 601

<210> 31039
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31039

agcttgtcca caaaaatagg ttnttgaagt ttgtcatttc aatttctcac taagtaaaat 60
ggatcatttt caaggtccaa cgcccttaaaa tgatcacttc ttaagtaaaa aagaatcact 120
tgataagaaa gaactacgta ggtctgattt tctcatccca attgaggaat acgtaggagc 180
aaagggaaac acccttgtcg accacaaaaa gagaaaaaat ataaaaaggg tataaaggat 240
ataaagacat aaaaagggaa cataaaaaat caaagtcacg tttgcacatt cgattaaagg 300
ttgccgtccc ttgggacgga cgtgtggagt gctaatacct tccccgtgcg taaatacaac 360
tcccgaacct ttcacttaaa agtt 384

<210> 31040
<211> 366
<212> DNA
<213> Glycine max

<400> 31040

tgccacccag ctgcccagg cgagcagggt tgcttctctc agaagcaaca gccttctgga 60
ggaatcttcc ggagggccca agtgggctg gttgctattt gcacccccat ttttactaag 120
tacacccctt gccttttttt ggtgattctt ttttggtaaa gttacggaaa cttacgaatt 180
tcgtaacgat acttgttttt tttccataat gttacggaac cttgaggatt acataatcat 240
cccctttttg acttacggaa tgttacgaaa cctcactaat tgtgcaacga tgcttccatt 300
tgatttccgg tgtgtcacgg aaccttacgg attgtgcac aatattttct tttgttttcc 360

ggcatg

366

<210> 31041
<211> 381
<212> DNA
<213> Glycine max

<400> 31041

agcttttgtgt ggagcttcaa tggatgaatga gggaggaaga aaagcaacgt gagggagagg 60
gagagagagc ttctgaaaat gtggggctga gtgaggagag agaggggtgc tttttggttt 120
aaataaaagg gttttctctt tttctattat tttatttaag caatgccaca tgtctccatt 180
tgagtggagc aagaagggcc cactttctct ttttgactgt gacctatatt cagtcacaaa 240
agtgaagaaa atctgacctt tgaaacgcta aaatcctgcc tcggtttgcg tgccgtttct 300
ttgattccag tttctcgcgt ttctctgcgt ccgccggggc cagttttcga aagcaagcaa 360
tatatatatc aaaacgctca g 381

<210> 31042
<211> 439
<212> DNA
<213> Glycine max

<400> 31042

tataaaactc agctttacat ggatgtccga ttccggtgaca taatatatcg agacgctcga 60
aatcgaacaa cggaagctct cgataaattc gaatggatcat aacatttcac tcggatgtcc 120
gattcgggga cataatatat cgagacactc gaaattgaac aacggaagct ctcatgatat 180
tcgaatgctc ataacatttc acacggatgt ccgattcggg gacataactt atctagacgc 240
tcgaaattga acaacggaag ctctcgagaa attcgaatgg tcataagatt tcacacgaat 300
gttcgattcg gggacataat atatcgatac gctcgaaatt gaacaaccga agctctctag 360
aaattcgaat ggtcataaca tttcactcgg atgttcgaat cggggacata atatatcgag 420
acgctcgaaa ttgaacaac 439

<210> 31043
<211> 159
<212> DNA
<213> Glycine max

<400> 31043
 ataaacatat atagctcata tatatatctt tctcgcatga ggaacactgg ctctaatacct 60
 cacttggcta tcttgaagat cggccccctt tgccatgtct gattgctcta tcaccataac 120
 tgctgtctat gaagcccata gtcttcaaat ggactcgaa 159

<210> 31044
 <211> 325
 <212> DNA
 <213> Glycine max

<400> 31044
 gatgatactg ctaactctat aattataaat catgcttttg tattctaata tatttcactt 60
 cctgtatgct gcgcaaaatc tcattcttac tgggtgtcaag tttcagacct tgcgatgata 120
 tgggactgtg gaactatggg atatatccgg tagctttgtg ttttcagaaa atgatgttgg 180
 gaaaatcatg gcagcaactt ctgttagtaa ctgcaagagc tcacaatgca gtgggtgtac 240
 aaagcttgac tactcagctg actaatgagc acctttccaa ttttcaagta cctacttctt 300
 ctgtctaate ttccttttct tttaa 325

<210> 31045
 <211> 435
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31045

agctcacatt ttncctttatt ctgnnncttt nggagctgnc aaatgattgg ttgtaatacct 60
 tttatgtagt tatgtactat gcataatgcc aaaggacaag tcataactatt cagttttcaa 120
 aaggaataac cttaaactgt catcctatat tgcattgngg tgggggtggtt aagtaggaaa 180
 gagaaacata ataaatacaa aaatatgata aagggatata atgaaataaa aaatgttaat 240
 acacattntt atgtatTTTT attattgatt aaaatttatt anaacgttag agattctatn 300
 tattgttaaa tgtatntaac tcataattct attattntta anaagtttta attaacaata 360
 aagaatattt taaaataata tatggatctt tnttcacaat aacaacaatg aaattcanac 420
 ttaanatttc atgct 435

<210> 31046
 <211> 414
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31046

ttgaaaatat aatatcttga tttctaaaat acccattgtc tctccctctt tgtaaaccatc 60
 aaaaaggcca aagtgcgcaa aacatgaata atttaatcat acacaaagca taattttgtaa 120
 aacaaacata taagattctg atacatacat aaagaaaaac atgaataaaa ccaaattgaa 180
 atgcaaacca cttagtcata taacacacac cataaatatc atgttcagtc atactaagca 240
 aatattaaaa gaaatactaa gttttcaa atgttcagtc atactaagca 240
 gaaaacaaaa tactaataat aatagtaatg tctaaactga tagtggtggt ggaggggaaat 360
 taatgtagtc acgaatgatg gtgaaatctt cttcaacctt tgtgacatcctt gagt 414

<210> 31047
 <211> 262
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31047

ttacagcaga tactagtagt gaccactaa cctacaatta acatttctga atgtccttaa 60
 cctangggat tagaactaac ttaatggctg aatgatactg aaattgctgg cgaccaaagc 120
 tcacccctt cagcaacctg taggcacat ttggtctccc taaatgctga tgcctacgtt 180
 gccaatgag cccttaatac aacttgaact aatgcccttg tagttgatta acccataaca 240
 tacttttggt cagccaactt ta 262

<210> 31048
 <211> 159
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31048

ctgcaagctg atctgcctct tgtaaaagta tgaccttga attctcggag cttcgttgtc 60
 aatttcagcg tctgatatgt gaacncctga atcaacatcc gtgtgaaagt atgaccattg 120

aattctcaaa gcttcttggt caattccaca tctcacata

159

<210> 31049
<211> 377
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31049

tcctcggtgc catttcctac gaaggcaaac attggaaagt agttttacca agaaatgcta 60
ctcttaaaac aaaaatggca tacaacctnc tncaataaac acaaacatcg atgtaaattt 120
aaaagcaact tatgcacata cttttttacc aacgggtcact tgcaccagac atcttataac 180
taaaaaaaat gcacccatgt acaatcaagg cacctttcgt acctagatta ttcatatgta 240
cttgccaagt gtatntgcta cctacatcac atgcactttc tttgctaaaa tacatacatg 300
catactcaaa gcatttgggg taccaaaaat gcacatgtgc acattccgta tttctaatac 360
ttatgcatat acaaact 377

<210> 31050
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31050

agctttgtgt tgatttctga ttttcatgca aaatctanat tagtagatga aaaaccagtc 60
tagaagggtta atctaatttc agatcatgtg gtaaaatctg ccataacaac tntggggaca 120
ggagtcaatc ccttttcgaa tactatatca caaattaaag ttgctcaagt aaaaaatgat 180
acatatacca agccagtagt tagctctttg agaaaaccat ccatttctac caatttacct 240
tctgacccaa aggttcattc ttatattcct tatgtatcta ccatagttag ttggaactag 300
anaaagagca aattgtcacc agcaatgact actattgatc atgatctaac ataatatata 360
gtagcacagt gtanaacacc aaagaggata tcatgcttgc caccgaggac tacatacatg 420
actcacnaac atttannaat tcagaaacat tgacaataat 460

<210> 31051
<211> 418
<212> DNA

aataatagc

249

<210> 31054
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31054

tggaacanat atattgagct cttgggtcccc ttagagattg tgtaaataatg tctcctactt 60
tatcattaga actaacgaat tcagtaataa cttccttaga aagaactttc tcctggacaa 120
aatgacaatc aatctcaata tgtttaattc tctcatggaa tactggatta aaagctatat 180
gtacggctgc ctgattatca cagcatagct tcatttggtg agtatctcca aacttcaatt 240
cttcgaagaa gttgtttaat ccanatgagc tcacaagtgg ctacagccat agctctatat 300
tcagcctctg cactagacct ttgcacaaca tttgcttctt actctttcat gagacaagaa 360
tttctccaac agacacacaa tatgttgaag tggacgcta tcnatgggtga tcctgccatc 420
tgcttgcaaa tccactatt 439

<210> 31055
<211> 342
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31055

tccgtgcgag atacatttct ttatgaatac attatttcta aaatcccaac agtgagaatg 60
tgcaaaaatg acttccacag gtggtgccca aatttcatga gaatccaacg gttaacgagt 120
ctacgatcgt aattctacta agacaagttt gggatatgc ggaaaagaga gaggttttgg 180
gagaagaaga agaaagaatg aacttgcgag gagcananag catagagacg tatectaaat 240
gtaaaactga cctagtatgt ctctatttat agttagggtg ctcttagcct attatttact 300
ttattatttt ttacaaaaca tacttctatc ttactttttc at 342

<210> 31056
<211> 338
<212> DNA
<213> Glycine max

<400> 31056

taacaagatg agttgccaac agagagagtc aatgaataat tacctatcga aataaatttc 60

cttcttttct ttaaagacga tgatttgtct tcgcacacca caagatgctt ttgctttcaa 120

agagaatctc cataagcctt aatagtcact ctcaaggggc cgctgaaaat tgctaattag 180

atagcattat ttatcaaaat acatgtaatt aactatgagt tacataaatt tctagtcatt 240

taattttttt caacattaat ttctctttct ttatgatccc ttggccatcc caatttttta 300

agggaggatt gctttccaca cctggggaaa aaaaaagg 338

<210> 31057

<211> 369

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31057

agctttaaga taagtgtgaa aganaagata gcagcctcaa tgnngtgaa actggagtcc 60

ttgcatatga cagagtcctt tgcaaacggc ctatgcttaa agcaacaact gtacaccttc 120

aagatgacag aatcaagaat agtcactgag caatcggccg atttcaatta gatccttgat 180

gatttggaag atatggaagt aaagctggaa gatgaggata aagctctttt gcttttgaat 240

tccttaccaa aatcctttga acatttcaag gattcaatta tctatggcaa agatcaagac 300

attaccctan aagaagtcca tgcctcaata aggaccaagg agatgcaaaa acagcaagac 360

tcccaatct 369

<210> 31058

<211> 352

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31058

tgcaacaaca gtaaattctat tnttgtaaa aatgaagtaa ctaactgtca caacctaccc 60

ttcggcgagg gggagacgca tgactcgagg gtgcgtgttc caagaaagat atacgcgcgg 120

agtcgccacc aacgttcatt taaggaaaat gtcggaaaaa ccggaaaaga cgtgatctac 180

aaactctaag tgaaagggtc gggagttgta ttacgcacg gtgaagggtat tagcacccca 240

cgcatccgtc acaagagacg gtaacctcta atcaaattgtg caaatatgac ttcaattata 300
 tttatttccc tttctacgtt cttatgtctg tttattcctt ttatgtatta tc 352

<210> 31059
 <211> 486
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31059

gacgagatga acattgtgga ccctttgatc acgcgacaca tanataactca gccttgacta 60
 acagagtcgn cncntccttg ttttctntga acttggttcta tatctcacct gtactccctc 120
 tattgccacc gtacttgga cagggtatg ttgcttacta cgatgtggag cttgatcctc 180
 aatatcttcc aatctatccc caggatcggg tgtaaacact actcataccc attccaaatc 240
 cctgaaatgt cctgaacctg atgcaacaaa aacacactcc ccatgaatcg aaaccaacg 300
 atcactgcca ccgtgtacat ccgccaatta aatgttcttc gtgctgacct ttaactgcaa 360
 tcccacatca caatgtcaac ctgacaattg tgatcttggc taaaaatcat gagccgtcgc 420
 ctgttcaata gaacgacctt cgaacctgac atctatcgag cctatctgaa aactctgcgc 480
 tgcact 486

<210> 31060
 <211> 294
 <212> DNA
 <213> Glycine max

<400> 31060

cttcatcaat ggagtccttt gcttcttgaa gatcaatgac agtggaatgc aaaaggagga 60
 aaggtgattg gagatgccac ttcaaggaga agagagtcaa gaacaagttc accaccatat 120
 gaagccatgg ataagagctt gaaagttgga gaaaatgagt ggaggagag ggagagaatg 180
 ggcacgaaat ttatgcctcg aatgaagtct aaaatttgaa gtgtaatttc tcaaatgatc 240
 aaagtagaaa taatgcacac aaaaagcctc tatttatagc ctaagtgtca catg 294

<210> 31061
 <211> 490
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31061

tatgaaccct gttganaccc ttgttgaaac cattggatan cctctcnana tnggcacnt 60
tggtggggat cttggtgccg attattgtag aggctttatg acctatctta agatttgaca 120
ctgacgattt cgaattttac tttcctgaac atagcgttgt acatgctgtt tcggcaccaa 180
gaccactgg gataagtcgc tcatgggaca cgggatctaa gtccttttgt taggtctgcc 240
tgagttttac tgctgactc ttttctttca agatattctc ggtcttaatc tagtcaaagt 300
gcctgttacc acatgaactg acccttgagt acaccattg ttatgatatc cccacttgag 360
ctatatacct ggcacacaca cctatatctc ttcactctca tggagaacga gccactgcta 420
cgacatcata atggttagat agactcccat atcgggtcaa ctggcatatg cattttctag 480
ccactcttcg 490

<210> 31062

<211> 237

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31062

tgctatttgc acccccatctt ttactaagta cccccctc tgcttgtttt tggtgattct 60
tttttcgtaa agttacggaa acttacgaat ttcgtaacga tacttgtttt ctttccgtaa 120
tgttacggaa ccttgccgat tacataatca tccccctttt gacttacgga atgttacgga 180
acctcactta attatgcaac gaatgcttca ttngatttcc ggtgtgtcac ggaaact 237

<210> 31063

<211> 128

<212> DNA

<213> Glycine max

<400> 31063

tgagcttatac tacacacact cttcatataa ctaagctcac ctcttgaga agcgtgcttg 60
agaagaatcc tgaagaagct tgagcttatac tacacacact ccctatctta gctaagctca 120
ccccatgc 128

<210> 31064
 <211> 426
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31064

 agcnnagant ttcaanttta tntacngaag cagnngagga ancttgggca ttatctttcc 60
 aagtgatata ttctttcttt cctttcttat ctttgaaagg tttcttgta gaattttcca 120
 tttttcttct aaagatagga caatcaactc tcaggtgcct aggttgatcg cattcatagc 180
 attntggaac tgaggaggaa tcttctccct tcttctttgg attgaggttc aatctcattc 240
 gatttctttt gttctcaga annatattta atccttttac aaagagactg anatcatcat 300
 cttcttctaa attatttttt tcattcaagt cttctttgcc actttcttca tgaatagaag 360
 atgaggttnt gaatganatt cctttcttct ttcnttcatt ctcttcatgt tgggtgagtt 420
 cataag 426

<210> 31065
 <211> 432
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31065

 tgcattgattt acatctccct ctntctcaag caaattcttc ttgatatcat caaatcttc 60
 atgatttatt aaagaatttg aaaataaaga ttttgaatta ttttgtgatg aacatggtat 120
 tgaacataat ttttctgcac caagaactcc tcaacaaaat ggagttgttg agaggaaaaa 180
 taggtcattg gaagaaattg caagaacttt attaaatgat acttctcttn caagtatttt 240
 tgggctgaag ctgtcaatac tgcattgttac atcatgaata gagccttgat aagacctatt 300
 ntaaagaana ccccatatga gttatttaac ggtagaaaac ctaatatctc tcatctacat 360
 gtttttgggt gcaaagtgc tgtacttaat aatggtaaag ataatctang aaaattcgat 420
 gcanaatctg at 432

<210> 31066
 <211> 418
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31066

agccttnctg ataccagtta ccacttcacg canagaanac gaaaggaaga agaacgcgac 60
ttangtgagg tcaggggtga ggaaggagac cgaaaccact tcacgcaagc aaaaagaaaa 120
aaaatggtga gggatcacga ggaagaagaa ggccaacgcg ggagggaggg aaggagagag 180
atgaaccatt tattttttaa ataaaaaaaa ttaagccagg tgtacaaagg tatttttgcg 240
tcaactgttg agtgcaccaa caaaaatggt ggggtgcacct agcagcactc gccagtgtac 300
aaacatgaga ccaacatana ggatatccag ttcacgagtn caacatccaa gttctctttg 360
ttgggtccga gtgttatgcc ctagtgccca aaaaactntc caatatctca tatactcc 418

<210> 31067

<211> 371

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31067

catttatctc atccattca acctaaacat ttcataaaag tgaacatcat aatcaaagct 60
tanatattca aactaggaag aaaaattatt caattcaaga ttaagaaaat tctctaggat 120
aaaaatcatt ntatgaagg acatatcana gcaaaacatg agtgcattga ccacaaagtt 180
gaaagaattg acaccataga tttagttatc catattccac aataagttgc ctgggtcana 240
aagcatttca agacacaatt agccaaagaa atttaattatt ntgttgcaag aataattttt 300
taaataaaag tagctacagt acaagtttat gaacatctat cacaacttat accaagaaat 360
tcttgataat g 371

<210> 31068

<211> 429

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31068

agcttaaggt ttngnattct attatananc acttcatcac ttgcttggtg aagatcatca 60
agccggtgcc atgcaatttc ctgcagggtta cacaatttct cagtattttg ataacctggc 120

agtggaatc acatatagag taatttacca tgatctttaa cataagaaca aagaaaagca 180
 atcatgttga aagttctcag ccacaacaca ctcaactgga ggtgcaagga aaaaacagta 240
 aagaccanac atagatccaa aattcagagt aaaaaagatt caaattagtg gaaactctgg 300
 ttttccttta gtttctctgt nttttgaaat tgacttatca tcccaggatt gtacttttac 360
 attctactct aaagaagata tcctagacta aactactata tangagataa ttaaagataa 420
 ctcttagta 429

<210> 31069
 <211> 413
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31069

tgtcaaagaa tccaacctct catggtagaa gcaaacacat agaaacaaga tttcactatc 60
 ttagggatca agtgaacaaa gagaaactga aagtggagta ctgctacaca tttgatcaac 120
 ttgctgatat ttttaacaaa cccctcaaag gggagagggt taaaaatgta aggggcataa 180
 ttggcttgat gaacttanga gatcagaata agggagggtg tgagagttaa attnttgttt 240
 gtgtggggta gaattgtttg tgctttgaat ataagagaga gtaacagaat ttttaaattc 300
 ttgtataagt actagcctaa gtgtgagngg ttattttactc tgttttgctt gtataaangg 360
 catacatata tcttaataaa gaggatttat tcattctatc attttcagtc tct 413

<210> 31070
 <211> 402
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31070

agctntgagt tanaatttga ctcaccatan accttgaccc agcgtgagaa tgccaatcct 60
 taccctcgga agcaaaaaaa gaatagaggg gaaatttcca atcacagaan aagagaagga 120
 aaatttccaa tgaaagcaaa aaagacatga aggaaaattc cccaatcata gagtnggaga 180
 aagcaanaa aggataagaa ggaaaattcc ccaatcaaag agtgggagaa agcaaataga 240
 tgagaaagga anattcccaa tcanagaatg ggagaaagta aaaaaggaag aagaagaatg 300

acagaaagct cctgatcaag gatcgaaaga aaccagaaga aatgtgcaga gaggtctttg 360
gaccagacaa tatctgaaca gtacagaatt gtcaccaa ga 402

<210> 31071
<211> 357
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31071

tgtaaattatt tattggtata atttgccctgt tccattagge tcttaatgtc tttagagggtt 60
acttcctcgt taacatcttt tgtcttgaat ggaattgccca tgacagggtt attggtactg 120
tctttgatat ttggtagttg atatttgtgt gcgggaagta attccgattg gattaactca 180
ccatccttca cttgccaatt tgctatgaca ttttgtgttg aatcacctat gatgtcttgt 240
ttccaagggt aatctatatc ctttctgatg gcataagcat gaaaccaatc aaagaanaag 300
acattaattt tgactctttt cgacaaatcg tagaacttgt cttggatttg ttctctg 357

<210> 31072
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31072

cctacttgac taggcgacaa tggtgaaaca gttctcacag actcagactc gacagggat 60
gtgaaagaca atgtggcatt ccagagcagt ccattattac tgtatgatca tgacttcagc 120
ccttacataa tctgagctca tggtttgctc ttgtcactga tgacggagaa gtcaccggac 180
cctaccctta ctgtgtcatg cctttgatta taggattgat gagatacagc agggcctggg 240
taatcacgctc cgattcatgt gctcgagctg gcacagaatc aagatatcag aggtaggctg 300
ccaccgccat tgctgcattg taacctttat tcagttgggc tactgccaag acatgtgccc 360
ttatcaacat tagaacgctg catattcgag attgttctct ctgcaccgt cgaatctgtt 420
gacactgaga agactgtgga taan 444

<210> 31073
<211> 444

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31073

 aatttgttat caatacctta aactaataac ttatgacatt acaacatgca aagttggaac 60
 gttccaactc atggcaaccg taaaagagca aaaagcaact caaagatgcg ttgattgtca 120
 acagaattaa tggcaaaagc aaatttgagt ttggtagtga ctccttgctt tggcaciaat 180
 tttatcagga tagtccttcc aagaaagata tgaatcaggg caatccttatg gtcttttttgc 240
 atgattcatc tgtttcgcac aatgaacatc accgtatatt taatgtatat ggagaaatca 300
 aagaggtgag tttggcttac caattgtttg ttgcacaaag attgatctga ataatttatg 360
 tgtggntgac taactgtctt gagtttctgt tcatcaattt ctttgggtaca ctttnntgtg 420
 taatctttgc tggaggatga caca 444

<210> 31074
 <211> 303
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31074

 tgaaattgat caacggaagc tctcgagata ttcanatggt catatctttt gacaagaagg 60
 tcagattcag gcacataata tatcgagacg ctcgaaatta aatagcggaa gctgtcgaga 120
 tattcanatg ctcattactt ttcaactcgga ggtccgagtc gagcgcataa tatatcgaga 180
 tgctcgaaat tgaacaacg aagctctcga gaaattcaca tggtcataac tgttgacacg 240
 gaggtcagct tcacgcgcat aatatattga gacgctcgat attgaacaac agaagctctc 300
 gag 303

<210> 31075
 <211> 517
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31075

 tgatgttcat tgcattcngg cgaatcactc gacccgggat ctgtgagtcg actgaggcat 60

gtgaaacaat ttctctctaa gaattttgac atgaaggata tgggtgatgt atcttatgtc 240
 attgacatta atattcatag agataaacct cgaggattg taggtctatc acatgaaatc 300
 tatattaaca acaattttaga gagatttang atganagaat gctcaccaag tgcgctccc 360
 att 363

<210> 31078
 <211> 400
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31078

ggacctatga aactcagctt tacatatttn ttccaggaga tagttgtaaa aatatntatg 60
 caaaagggtga taattgncca tgtattgcat tgctcttagc tcaacattca tccaatgagg 120
 ggtattgtga gaagagtga aaaacgcggt tttgtagatt aaaacaaatg ccattggagc 180
 taacgtggaa agacanagaa attaacaatt gcatataaaa aggggggtttc tgggtggtaga 240
 caatattgta agagaatagt gttggaggaa aataccttaa tttgaagtaa acatgggtatc 300
 caacctgtgg ccaactcgat gcttttttga ggaatgtgct ctcgctgctc agctgcaact 360
 gtgccttact attactaaca ggtcaatttg atgatggaca 400

<210> 31079
 <211> 380
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31079

agctttgaat ttactattca atggagttga caagaacatc ttcagactga tcaacacttg 60
 cacagtggcc anagatgcat gggagatcct gaaaatcact catgaaggaa cctccaaagt 120
 gaagatttcc agattgcaac tcttggttac aaaattcgaa catctgaaga tgaaggagga 180
 agagtgtatt catgacttcc acatgaacat tcttgacatt gccaatgctt gcactgcctt 240
 gngagagagg ataacagatg aaaagctggt gagaaagatc ctcagatcct tgcctaagag 300
 atttgacatg anagtcactg caatagagga ggccaagac attngcaaca tgagagtaga 360
 tgaactcatt gggtctcttc 380

<210> 31080
 <211> 409
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31080

agcttatgag ttatctaatac aagattcttg attgctctta aagttgtaac atgtgggtctg 60
 actagtctcc actttctagg taaacttcgt aagattttat caacatgatac ataattatca 120
 taatgtatac ctagagagcg gagctcggtc agaattgattt ggaagtttcc aaacatgggtt 180
 tgaatatctt ctccctcttc catattaaag agttcatact tatgtgtcag aaggctcaac 240
 ttgttacgtt ntacgttaga ggacccttcg taggtaattgg ataaggtatc ccacatttgt 300
 ttggcgcttt tgaagttgtg aactttggaa tattcttgcg cgtagttga taagaaatat 360
 agacttatga tcatccatcc atttgcctt ggggatcttg ttcttctga 409

<210> 31081
 <211> 320
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31081

tccttgagag gcttctttga gaagctagag tcttaactat ccacaccct ctaataacta 60
 aactaacctc cttgaaaata aaacatggat aaaataacac aacaaataaa atcaaacatc 120
 aattataatt gctaataata tttcaagggtg ttacagcttg tccaaagtag ccttgggcat 180
 gatgttgagg gaagagccat tgtcgataag cactttggcc actatgtggt gatggaagct 240
 tgcttggtga gcttctatgg aggctggatc ttgagcttc aatgaggtcc ttcaatggtg 300
 attntacacc atggagatgc 320

<210> 31082
 <211> 373
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31082

taagcttatt cttttgacag nntattatca tgcacaacct gcaagaagtg gctcataaca 60

ggccaatcat aactatggag cattttcttg agcaagtagc ctggcctgaa gctcaacttc 120
 cattggtgag acccaacgag gctactccgc ctgagccac ctgtgcaggt tgatccagag 180
 ccaactaacc cacaatctct agtggtaaat ccactatctt ctcttgagcg tgaagtagtt 240
 ccccatctc cacctctgat tatcatctcc gatgcatcat ctgatgaagc agctgcccc 300
 tctgatcacc anaaggagaa aacagctgac cttctacttc ccctagtgga ggaanttctg 360
 antcgtcatc tgg 373

<210> 31083
 <211> 393
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31083

tgtgacattn tcaattctac atataaccct gtaagttagc tatctattga tcttattttg 60
 tttatagggt tctaatttct atcatagctt gggtacatat atttatattn tattgatgtt 120
 gcagggtgcac atttgggact tttcaggaca gacaaccag ttcttcttga atgacaaatc 180
 tagattgcca aacgattctc ctggtcaact tggaaaagag gtagttggtg attgttaata 240
 gattggcacg tgtatcaatt ntatcacaag tagtaaagat taatatggaa gttcaagtat 300
 cgaatccacg aggactttgg ttgtacttta gtgattctaa cccaattatt aagcaatgag 360
 aagaagtaga agagaaaatg aattgtaagt gtg 393

<210> 31084
 <211> 435
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31084

ttagcttatt actatntcat gnnactatct atgattttgt tacttttctt tttcatgatg 60
 aagttattat catgtaaact gaacaactct caacatatct ataatgcatt aagttatttt 120
 gtagcaagat ttaaacggtt ttggtgtgtt tagtgacaca ccagaacatg acattgctaa 180
 tcattaaagg aaatctcttc taaattgagt aactcactct agagggtaaa gtgagaaatc 240
 atagttgttg atgaanaaat caacaatcag ttaggtgcac atatatgaca aatcatggat 300

cttgtggaat attgaannat gaatttttagt anatggtcta atttatatattt ggttcttaat 360
 agaaatttgg tntatgaaga atctttaata aaataataat ttttttttat tcttgacact 420
 tatntctagt tccta 435

<210> 31085
 <211> 310
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31085

ttcttcgctc ggtgctcctc tcatggggta ccctagttgt cttatagcga gcgcgggatt 60
 gtagttaata caaccctcg ttcctaccag cggaatgttt ggtatcctc cacatgagaa 120
 gaggaccctt tcttttcctt ctttccatcg ggggaaccâa ctgatngttc taccttctat 180
 cccggccaag agctggtccc aatctattct cctcttttca gtacacgagc gatggctcag 240
 gagccgacat ggatgtcttg ggtcttggtg gaacaagtgc gaaaccaacc atacacagag 300
 ggcgggtaag 310

<210> 31086
 <211> 191
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31086

ggattcattg gcctcttacc tgttcatttt atgtgtggaa ggcctatncc ctatgatcaa 60
 gaaggtggag agtaaaggac aggtgcctgg ttttgggttt gtaggaatgc ctttctatca 120
 gtcattcttg gtttgccggg gatattctatt ctttgagcat ccaataagga gtgtatgagt 180
 attcaacata t 191

<210> 31087
 <211> 175
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31087

ttcctttgac tcgattagct aatgatgtga actgcacgta atatgactga actgaaccaa 60
 tttgagttagg ttttaacaac tgagatctan gacattcaca cggatgatggg cccaattctg 120
 tctctaggct tgcgtanaac aatccatggt gtgatgattt tcacgagtat tattt 175

<210> 31088
 <211> 442
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31088

gcttgctgat tacattctcc cncctttctca agaaaattct taattcttct tgacatcatc 60
 aaaatcttca tgatttaca atacaaccca naaataaaat aaaataaaac tggacgacaa 120
 ataaaattgt ttgctctttt caagtccaag ccggttcagc ccaattctgg atccaagccc 180
 aattgcttat aattctcttg aaattaaatt aaaacacaaa attagtcaag taggtccaaa 240
 tgataaaaact gcataattaa tttgacaatt aagggttaatc agtaattaaa atgggtgacag 300
 aaaggggtaa gaaataggag aaaataatga cacatcaata ggcaacttcc cccctatgg 360
 tgattagctt gagtctcaag gaagtttcan accgagtggc atgcccccaa gtacaaatat 420
 ttttctcat gaaaaactac ta 442

<210> 31089
 <211> 377
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31089

agctngtact taacttatga naaatcaaga acaaacttgt tcgcacatcg ttcgctgtga 60
 tgatatccac tcgacaaggt ttgaagtgga ggagaccttc aatcctataa cgcaacgtgg 120
 cggacaaaaa tgggcagtta actngaattgg ccattattgt caacgcggaa ggtatnttgc 180
 gcttcactat ccattgtcac acattattgc agcttgtggg tacgtgagca tgaactacta 240
 ccaatatata gatgttgttt acaccaatga gcacatctta naagcactat ccgcacagt 300
 gtggcctctt gggaatgaag cggcaattcc tccttctgat gaggcattga cactaatccc 360
 tgaccaact acaattc 377

<210> 31090
 <211> 388
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31090

ttacgaagct ntctgggcct tgcgggcttt tataggagat ttatcaaggg gtatgcttcg 60
 atagcttccc cattggtggc agcaaccacg gtggagcctt tccagtggac cgcggcggct 120
 cagctcgcat ttgacctctt gaagaaagcc ttgttcgaaa ccccggtact tgccttgccg 180
 aatttctagc taccatttac agtcgagacc aatgcttctg ggggtgggcat ggggtgcaatc 240
 ctctcttagc agggccacac aattgcatat tttagcaagc catttttgcc taagcttcaa 300
 cgatcgtcca cttatgtccg agaattgttc gcagtgatgg cggcgggtcaa gaaatgggtgg 360
 caatacctcc tcggtaccg gttcatca 388

<210> 31091
 <211> 445
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31091

tctaatttac tacattcatg caagacttaa ttggtttcct ttnttaacaa attgaacatt 60
 ttgtaaagac aagggtctaa ttataattga aaaaaagaag attgtgttta attaattaat 120
 taccaaaggt gaaggagctg aacattatgc ctaaagaatc agagagctct cggtaactct 180
 tgtacatctt caagtccact ttgcgaaggt aagggtgctc atccatgcta actttgacaa 240
 agcttgcatc agggctgctg ttcttctcgc tctcttctcc aacgctcttt tgcacagcca 300
 acatgttctt ccggaaggac cgcacagggt gccaacccac cacctgcgtc ctacacattt 360
 caactcatca atatcactct atatattatg atcaattaat acgcatcatg aacatatatg 420
 gcaatcatat aacgaagtta aaata 445

<210> 31092
 <211> 273
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 31092

ttaatttttt tgctttacct tctcttccat tgnntgttct tcattttttt ctccatgtat 60
ctctcacat gtcttgtgct aaatgttggt aacacgattc tatagagttt ccaccgatta 120
aacttgctat agaagctaga attgattntc tatggttcaa atttcttggt attgttcttg 180
aaccatgaat tgtgttgagt ttaagttgct ttgagttttg tcttggtatt ttttgtggct 240
gaaacctaaa ccataaaatt cttacaaaat att 273

<210> 31093
<211> 334
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31093

tgatgtcatt caaaacacac tatgtagacc taaatgaaga ctaatcattg tttatttaat 60
tggattcatt atacgatata atttgttgta acccgttact aaccaattaa tattatcaac 120
tactcgtttg gttaagcaag gaaattgttg gtccaacaaa aatcatttac gcgtgcagca 180
tacatcattg tcataattga caacacataa tgacatgcat gtgtattaca gtttgagcgt 240
gacaacacat tggctgactt cagtacacat tntgaaacta gcagtcgctc gacaacacat 300
tggttgactt gactacacat tagcgacaac acat 334

<210> 31094
<211> 520
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31094

catgactatg cgatgcaata ctagccannc actcgaccgg ngatccttna gtcacctgcn 60
gcatgcaagt ttggcttcat tatggagcag agaaactcca ccagtaacn ggggaccacc 120
ncgacaggaa agctctttac catctcgac ccgagcctnc agcttatcta tctcctctgt 180
ggagctatcc agggctctgg tcaactggct aatcctctcg actcgacaag atagagagtc 240
gaacgcttct ctgcttcttt cgatgggtgg tcggaactcg taaccgcttg caatgatatc 300
aacagcccct ttcaacaact ctcccgcagc agctccagcg aacggctctg ccatactac 360

tcacacacac tgcgaggaat gaaatgaaag ctcacgaata atatgtacta ctcacacgac 420
acaçcaaacc gcgcgttttt tcgttgacga tgactacgct atgatctctg agctcgcgta 480
atcanaacat agaaagacac tgtattctct tcgattaccg 520

<210> 31095
<211> 347
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31095

tatgcgcata tttccttaca aacgttctct tgcacaagac attctattaa ccgaataaaa 60
tgcaccata tacaatcaag gcagcttcgt tacctagatt atttacacgt acttccaagg 120
tgtatttggt acttacatca cacacatctc cttggctaaa ttcacataca tgcatactca 180
aagcattntg ggggaccaa aattgcacat gtgcacatct tggattttct aatacctata 240
catacacaaa cctcatgatg aatcttgact atctacacaa taagggtgcta catttcatgc 300
tcttttcaag tttttgctac ctaaggccgc atgcaaattc aagtata 347

<210> 31096
<211> 302
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31096

tgtctatcgc acttacaatg acaaacatca tcttttagcac aaactatgag caccaaccaa 60
gaaatgaata tagcaatgag aaaacctata gaatccaccc ctgtcctgtg tcctatgctg 120
acttgctccc atatctactt gataattcaa tggtagccac aacctctacc aaggttcac 180
aacctttatt tttccgaaaa tacgactcga acgcaacgtg tgcttgtcac ggagaagccc 240
cgggtgtgtac cattgagcat tgtanggctc tgaaatgtaa ggtgccaggc catattgatg 300
ct 302

<210> 31097
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 31097

tgcttctaca tgtttttcta tctattacat tcgtagntag aaatattttt gttctataca 60
 cacatgacac ctacaccttt gcacacatgt tgagatatta agccctatac ccgggtctgt 120
 gtgagacata nggagtggag gttgatctat ggtcatgttg ggtcttcgac ttgcttgata 180
 acagtgatgc ctcatctaga gttttcttct ttttgctgat gcattgtcac tggtagatcc 240
 taccgccaca atgttggttac ctaagaggat gatattctta gaagccaatg agttacatga 300
 taccaccttg ggagttgcac tagaaggagc tttggatcct ttcatangtc ctgaatatga 360
 cacatacaac tcactt 376

<210> 31098
 <211> 428
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31098

agtttataat tattaatttg acggttctcc acattntggt tcttgaaatt ggtattcttt 60
 ctatgggtga gaaacttgcc ccatttgtat ggccatatatt ttcaagggtt ttgctcattt 120
 atcagcttct ccaaaggctt gacctcagtt tcatgcgaaa ggtagaagaa tggaatggag 180
 aacctttctt tctcagagtt gaccaccact ctgtgttcca cactctcata tgcattcattg 240
 ctccaaacct gcacaaacaa ctccaatcct canataaaaa ttctctccac taattacggg 300
 aataaatagt tcataaaactc aaagattaga atatgttntt tttaagggtca tcatgacatt 360
 ggtgtgaagt atttatactt ttgttagtga ctaatctctc ctgngaagct gggcagacca 420
 ctattagt 428

<210> 31099
 <211> 403
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31099

tctacttatg tggcagggca ggcttccttc actttcttgt ctccaacgcg agctttgacc 60

actgttcttc cttcccgga tgcttctttt catgtccgcc tgagtgggct tatagcctaa 120
 accatacttc ccacgatttc cttgggtatc tatcaggcta gttatgccgc cgttggtttt 180
 gcctaaaccc atcccgggtt cataaccgtt ccccaacata actcgggcca tcattaccgc 240
 tgcacgggac agacaaggct gcccaaagag ggagtccacg gaggaatgc tgaccacctc 300
 anaagactgg aaagcagttt ctaacgattc ttctgaggct tccacataag gcatggagga 360
 tgggcagctt accaagatat cttctcggc tgacacgatg act 403

<210> 31100
 <211> 333
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31100

ctaaaaaatc ctatatttct aggggtaccct acctatatta tggagcccta aatacaaggc 60
 ccaaaaataa tgaaacctta atctaattt taaaaagat agtgggctcg tacttagccc 120
 atggggcccaa aatctaccct aagggtcata aaaaccctag ggccttctct tgcattctctg 180
 gcccaatcta cttggagttt ctatccaatg cccttgcggn gtaagattgc atcattccct 240
 cccctagaa gaggatttga cctcaaattc cgaggctctt gaactttggg ctttttttct 300
 cacactatan aagaacaaaa catatgtata gtg 333

<210> 31101
 <211> 156
 <212> DNA
 <213> Glycine max
 <400> 31101

gcagctctat ggtaaattgtt aatgtggtga agggaaattc cggcgtgtta aggtttcagc 60
 attgacggcg acgcagagaa gccgtcaacg tcgtccgaga tcgtgttgga acccatcata 120
 gacttctcgg gtaccatcac attgccaggg tacaag 156

<210> 31102
 <211> 322
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations

<400> 31102
 atatagggcc ctctgaataa acttttgcac aaacacttat atgagaagaa agaanaaaaa 60
 aagcttctca ttgttaaggt catttgaacc aaaattctca accaactctc ctatngaata 120
 aaatcaacgt atgcacttca aattttatac aagatattct tcatgtaact tctccaaatg 180
 tagattnnta attatgagaa aaacttaatt atttcatctt attttcttct ataagtactt 240
 attgaaaagt ttctccgaac atgacaatca ttaacattaa naactgcac ctacctaaat 300
 ccatntgcta gcaagatcat ta 322

<210> 31103
 <211> 364
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31103

taagcttacc tggaagccta ggaattaatg cttttatata taccattctc tcatagaatg 60
 caaagcatgg aaagagaaaag actgagtaga aaagactggg aaagaggcag tagaccctc 120
 aaattgttct tccttttttg ctttctcagt attcctcttt aactctaggt gctacagatc 180
 tctatttatt gccatactaa accaacaata tggaattaat ctgttttatt tctagtatcc 240
 catcaagcac caagtgaaaa aataatacca tcccaacata cagttgtact taccacctac 300
 accanagtaa tagaacctac acattaaaaa atattaatag tttaaaggat agtatttttt 360
 ttct 364

<210> 31104
 <211> 316
 <212> DNA
 <213> Glycine max

<400> 31104
 ttgagccaaa atcctaactc accataaacc ttgacccatg gtgatatatg tcaatcctta 60
 ccctcaggag caaaaaagaa gagaaggaaa atttccaatc aaagaaaaaa aaaagagaag 120
 gacaatttgc tatcaaagag aaagcaaata aaaaaagag agaaggaaaa tttccaatca 180
 aaggataaaa gaaaggaaat gaaattccca atcaaagagt gggagatagc gaacagaaaa 240
 gaaagaaaac tcccaaccaa agagtgggag aaagtaaaag gaaggaaaga aagctcctga 300

tcaaggatcg aaagaa

316

<210> 31105
<211> 414
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31105

agcttatgaa tttctatttg caatggtatc tcccaaagt gagtagcatc tctactatga 60
taccagttat gaaaacaatg cacgaattct aaaatgcaac acttagcaga aagggtcaat 120
tgtaaagttc atatagcaat tcaattctaa tccatatata acgtatttta tatatattca 180
tattcccaa gagtctactt ttcaaata attttatttt catcaaactg tatgtgaatc 240
aaacaaagta aaaaactatg tgaagtatgt caaagttgaa aattgaaaac agcatgtgtg 300
cacaaactnt caacaccaa taatttagaa atgactctaa gagcccatc tcatggagga 360
taacctccca naccanaatn gacattaaag aanatagaaa ctctcaatac cttg 414

<210> 31106
<211> 307
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31106

tggaaccaa catttaagat atgataagat tgtagttta aggtaatgat tgaaattaaa 60
tttaaaaata tattcgattt tatagttata aaaaatattg taacctacat ttaaatttag 120
actattatca gattgctagt gtaagataat gattgaaatc aaatttaaca atatattaca 180
tttggtagtt ataaaaaata ttgaaccaa aatttaagat ttanaatata tctattaatt 240
catatgttct aattntttta cgagtatggt ttagagnaa aaaattcatt taatttattt 300
acaaaat 307

<210> 31107
<211> 167
<212> DNA
<213> Glycine max

<400> 31107

actgctccat attactgata atcatgggac ccatacccca ccaaggtatc aacctcattc 60
 tccgaaatac actcaacgca cgtgtgcttg cttgacaacc ccgggcgttc attgacattg 120
 aaggcctaag cgtaagtcag gtcaattgtg cggctgctga attcaga 167

<210> 31108
 <211> 296
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31108

cttatccgga tgcagaatat attccgaaaa gactatctgg atgtaagaat tacctttaaa 60
 agttttccaa tataatttac attcaaatat ggaagaagta atgggagttg aatttcccat 120
 ttgaatgcga tggctggaag gcttccttta ctgggctgca agtttgcacc gaaggaacca 180
 ttgcttactg cacctctaaa ttactaccta cagccacat cattttaaaa caattaaaat 240
 tttcnaaagt naccgccgacg tgtccttcgc accccattcc ccgtcccatg gatgct 296

<210> 31109
 <211> 462
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31109

agcccanncg ttaggcctta attttgggaa caatcncng nattggcaaa gaagacatca 60
 tatgccaaagg gaacaatttc ctccatcac tggaggtata tacctagggt aagagcgagg 120
 ttgattcata tttctaaaaa tttgagacaa aagttgacct aatacgcttc tacaatcttg 180
 tcaagataag ttgcatcgag gatgatgaag tcgtccctat atacttgtaa ggtctcaata 240
 actatatata ccccgaaag aaaactactt ctttgacaaa gacggtgttg cactattaga 300
 aattacactt tcaacatcgg ttatttaggg cattctacat cggctctaan accgatgttg 360
 aaagtgatga tgttgaatgt atcatcggtt acatcggttt ttaaaaaccg atgttaacat 420
 anatatgata acatcggttt tctaaataat cgatgtaaac ac 462

<210> 31110
 <211> 384

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31110

 ctaagcttaa gctgctcaat tgctccaggt tgctgcatgg aagggcaaag gtctgtatgg 60
 tggtcagcag aagagtacaa accacanact cttgcgacag gtacagattt cggattcaaa 120
 gctagctggg ataccaagtt aaccaatgca tccaagttgc cttcaagctt cttagtctca 180
 gatgatgcag ntgagtttgt agctacctca tgcactcctc taatgactat agcatcattt 240
 cttgcgctaa actgctgnga gttggaagcc atcttctcaa ttaaatttct ggcttcagta 300
 ggagtcatgt ctncaagggc tccaccactt gcagcatcta tcatacttct ctccatatta 360
 ctgagtcctt cataaaaata ttgg 384

<210> 31111
 <211> 167
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31111

 agctangcac ttaggatgga atgatattng nagctaata acaactttct taactatttg 60
 tccttcagaa tttactgttt cttntgtcta atatgtaaat ataaattgta taaggctatg 120
 gtgtaaaaac atgggtctacc agctcaatat ctatgggtta tgcttct 167

<210> 31112
 <211> 403
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31112

 ccttgggcaa tccttaccat acaagacaat tttgtggtgt tgagttagac cctaaacccg 60
 aaatctaaga tggattaga gcctatccta gatacattgt tggggcacca acattgccac 120
 gctccaggcc catagcccta ggcattaggg ggtgtgttgg acagcttctt taattgcagt 180
 cactgctaac ctgctntaat tgcagcagca tatgagagtt ggttgccaat gtctcagaan 240
 aggctaccta tgaagggact gaccagaacg gctgagttaa gcgtcgtagt gtgcaatcaa 300

tgagtctgaa acatcaactc ttaggggggtt gagatccac attaactaga gataaggcct 360
tagtattgct tataaagttt gggcaattct caccatacaa ttc 403

<210> 31113
<211> 281
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31113

tatcatctcc ctcttcatca tttggggcgc tacttgagct gccagatccc tccatctttg 60
ggtgtattct ttgaaagatt catgctccat cttgcacatg ttttgcagct ggattctatt 120
cggagccata tcagaattgt actgatactg cctaatagaag gcaaccatta ngtcctttcc 180
gagaattgac tcatgaaggt tccagattag tataaccagct gacggtttcc ccagaaagac 240
tgtcctggaa gaagtacatn ncacaatttt catttttcga g 281

<210> 31114
<211> 251
<212> DNA
<213> Glycine max

<400> 31114

ctgatggcta tgagaagaaa tcacatgttt gtcacatca aaaaggggga gaatgtgaat 60
gtatgtatac atgattttga tgatgtcaaa agaagaatca aacaaggctc attttgcttc 120
aagattaata ccagattggt tcaacaaaca aagccttgat tcaagaattc ttcaagatca 180
agccttgctt cacaatgaaa ggtttcaagt cattcaaggc acatgtaatc gattaccaat 240
acatgtaatc g 251

<210> 31115
<211> 257
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31115

tggatccata gtctccacgc cacatctgta aatgaaggan tccaccctag ccaatgagat 60
tgcgaccatc aaggctgac aaaagcaagc acaataatgc tatgctgaga gccttgaagt 120

aacaccctat ccttccacta gggagctggc caagcctcac cctacagcga gtgaaggtag 180
 tcaagtcatt aacaaagggc ttacaatccg agccttcatt gtttaccaa caagcctgga 240
 cgatgaattt gatatag 257

<210> 31116
 <211> 386
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31116

accataagag ccttagatat tngttngatc aaaaagagct taacatgagg cagaggagat 60
 ggtagagtt cctttaagat tacgattttg agccttagcta tcatccaggt aaagccaatg 120
 tagtagctga tgccttaagt agaanatccc ttcaaagtgc tgctttgatg gttagagact 180
 tggatctctt anagcagttt agagacatga gtttggcatg tgagatcacc tctaatagca 240
 ttaagttggg tatgttgaga gtcaccagcg aactcttgag cgagattcgc gagggtcaga 300
 agtctgaccc attcttgta actcagttag agtccatagt cgcanggaga gagagtattt 360
 ttagagtggc tactgatgga gtcttg 386

<210> 31117
 <211> 344
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31117

tgtccatcaa acaactagga acagtatgga gtctaataca tcaattcata tttcattntt 60
 gcttatccac tttggacttt tctttntgtg gccttttaga gggaaatgca gaaaggattt 120
 tctgtgcac tagtgaagaa agaagcatat tctcaactca actaccaaac aaaaggata 180
 agagactgac cggtgagact tgagaagggt ctttctattg cattnttcct accctgaatt 240
 nttctatgat taccacgtta cctttgcctc tttgaagagc ccaagcctgg agagggaaaa 300
 tgtcatttaa cttaacccat gccttacatg atangtcaca aaat 344

<210> 31118
 <211> 452
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31118

ctttctatatt aatatagngc ngctactggc cttaggtgag gtaatttatt gaaaacacag 60
gtctatcaaa tntatgttta attggatgat atatatatat atatatatat atatatatat 120
atatacatat atatatatac atatatatat atatatatat acatatatct atatatatat 180
atatacacac acacacacac aacctttcat ttccacatat atacaccac agacactctc 240
tctcgcagac atatatctct tctgtctctg acacagtgtc tctctcaaac aatacacact 300
tctctcgcga gcatccttga gcaactcacc cggcgcgatgg agcgcgatgaa caaactaata 360
tacgcactgc acacacatta tcatcccacc gtgatacccc gccactacct tgtgagggat 420
cttctctcac tcaacagaca ctgaatacac cg 452

<210> 31119

<211> 389

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31119

tgctgatggg ggtgttgaac tttntacttt tgtttatcat agtaataata tgggggatgg 60
ggcacttgca tagattatcg atttgatatg agttattatc tacaaaagag agttctcttt 120
agtcttgggg ttgtggtggg gacattgctt tgcaatatgc atattcgcag ctgtggttgt 180
gtaatagtaa tagcattttt taaagcatct ttaatagcat ttttgtgtaa tagtaatagt 240
ttagttttta aaataaagta ataatcctta gaaaacattn tctattatcg taataaactt 300
ttggtagttt atttaaaatt aaatntatca ttntttacca tgatattatt acatcgatgt 360
ttataaagac cacattatgt atgaatgga 389

<210> 31120

<211> 406

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31120

nttgttttca atgacgagcg tctcgaaatc ctacgggaca ctattggaca tccgagtga 60

aagttattgt cgtttgaatt tgttttagagc ttatgttttc aattacgagc gttttgatat 120
cccacgggac acaatcggaa atccgagtta aaagttattg tcgttagaat tttctcatag 180
cttccgtttt caattacgag cgtctcgata tcttacggga cacaatcgaa catccgagtc 240
aaaagttatt gtcgtttgaa tttgctcaga gcttcagttt tcaattacga gcgtctggat 300
atattacaag actcaatcag acatccgagt taaaagttat tgcgttnga ctnttcatag 360
agcttctggt ntcaattaga gcgtcttcat atattacgag actata 406

<210> 31121
<211> 243
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31121

tgcattcgga attgcgaaag ccatgctcnc tcattangat tcgttctctgc catctcaaatt 60
aagcaaataca aacataataa gacaattata gtttctgttt gaataacctca cccactcaag 120
tgtatcacac aattatggct tttctctaata gaaacactct tgcctttttac cactctaatt 180
ccccttgagt tcttaagcaa ttcaagagat tatgtgccac aacaaagaac aattcaccaa 240
aat 243

<210> 31122
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31122

ttatcatttt actanattca aattgncata acatttcaact cagatgtctg attcggagac 60
ataatatatc gatatgctag aaattgaaca acggatgccc tcgggaaatt tgaatgggtca 120
taacgtttca caccgatgtc cgattcgggg acataatata tcgagatgct cgaaattgaa 180
cagcgggaagc tgtccagaaa ttcgaatggc cctaactttt cacacagaag accgattcgg 240
ggacataata tatcgagacg ctcgaaattg aacaacggaa gctctcgaca aagtcgaatg 300
gtcataaactt ttcacacgat gtccgattcg cagacataac tcacttaaac gtcctaaatt 360
gaacaacgga agcaatcgac aaatttgaat ggaataaca 399

<210> 31123
 <211> 400
 <212> DNA
 <213> Glycine max

<400> 31123

tgagatcttt agtcattct aaggcttatg agttattttc gagatattga gtattctctt 60
 gatgtatatg gaggagaaca caagctagtt atttatagag aaaataatta taatcgtctt 120
 taatcaatta aatctacaaa gtaattgatt aattcaacga agtaatcaat tagattatct 180
 ttttaatcga ttaaagtatt cttaccaaca tctggacata actcaagaac aatgtaattg 240
 attaaatact ccaagtaatc gattaaagtg ttcttattca cttctgaaca cctaagcgag 300
 agagacgtaa tcgattaaat cacttggtaa tcgattaaag tagagactcc tgataaatca 360
 gccactgtct caaacaatgg gtaatcaatt acgagatatc 400

<210> 31124
 <211> 525
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31124

cgattgtatg tanntcaatc gcanaccac ggacccggga ncctctaagn caccagcggc 60
 atgcagcnnn gangtangca ttagttgtag gagaacggag ggaggngaag cnaannnnng 120
 gagcgcacaa ncaggaacng gccangncan nagnnnngaa accatangcg ggacgaaaga 180
 ctaaagcnga aaaaacactt anggcngaca cgacgaagnc catggagaag ctaagaaaca 240
 ttctgctaac tctgaaggaa cacaatgtca atagttatac gaccattaaa cagatataca 300
 atgcacgaag tgcatttcgt tcgttcataa gaggaagcga tcttganatg caacatctga 360
 tgaagcttct tgaacgtgat cagtatatc attggcacag aatanaggat ggagacgtgg 420
 ttctgtatat cttttggtgt caccctgatg cagtgaatgt agtcaacgca tggatttcgg 480
 tattttgata gacaacacct acnaaacaga cctgtacaga ctccg 525

<210> 31125
 <211> 306
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31125

ntgaccattc gaatttcgag agtgcttccg ttgttcaagt tcgagcgtgt cgatatttta 60
tgtccacgaa tcagacatcc gagtgaaatg ttatgaccat tcgaatntgt cgagagcttc 120
cgttgttcaa tttcgagcgt ctcgatatat tatgtcccg aatcgaacat ctaagtgaaa 180
tgttatcacc attcgaattt ctcgatagct tctgtgttc aatttcgagc gtctagatga 240
gttatgtacc cgattcgaac atccgagtga aatggtatga ccattcgaat ttctcgagag 300
cttccg 306

<210> 31126

<211> 411

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31126

agcnngnnaa ttttctttat cccatnngaa ctntnngnc tttntcagta agttcatgat 60
cggcctagcc ttctcagcta tccaaggtag aaatcttate aaagaggcta tgcgtcctgt 120
gagtctttgt atctctttga aagtcttcgg actcctcctc tcaatgacga cttgacattt 180
atctagatta gcttgtatgc ctctttggga aagcataaaa ccaaaaaatt ntcctcctcc 240
aatcccaaga acacattttt agagggttaag tcgtatgtat gtttttggat ttgtgaaatg 300
atctcggcta ggtcctcaac atgggacttg actccatngg atntgaccac tatctcatca 360
acgtacacct ctatatttct acgaatnnta tctttgaaga tcttatccat g 411

<210> 31127

<211> 337

<212> DNA

<213> Glycine max

<400> 31127

tcttcaagaa aagattattc ttggtcttat atgattctat gaagataacc tacaactaaa 60
aatggtttct gacatggttc tgtaaagatg aaatttttga aaatggacaa gcaatgacta 120
tcgaacacga aagtaaactc ctgctttact ttttttattt gttatttgct tattttattc 180

tcaatttaga aataactcaa tggacaaaat aatttataaa aataacatat tagccaatga 240
 cctacattca atttaaataa atgggtcatga ttctttactg tcagtgactt ataacccaag 300
 ttaacaaaag ggtctattga cataatactt gtagttt 337

<210> 31128
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31128

agctncgata ttagctgttc cattttanca anaaacacaa gnggaagttt attcagaana 60
 ttagagctta tctcttttat cttagtgaga gtgattctcc taaattcttg agtgattcaa 120
 gaacaccctg gctgtatcaa aggactttca caacctttga gtgttgccct cgctggaaag 180
 agtgattctt tccttccat catctccacc cttgttcttt caaaccacaa ttccagaaaa 240
 tccacctctg ccataatta tctcgtggcc ataactccca ttntatgcac tcaaattaag 300
 tgattcttga gcctaaattg actntcanaa cgagaccttt cacctcgnrc tgaaatcacc 360
 tcattnggag cctgtagct tcagttattg ccatgtctat atttctgtcc agccaccact 420
 taacctacat gttaccatcc cattcatcca ttttat 456

<210> 31129
 <211> 353
 <212> DNA
 <213> Glycine max

<400> 31129

gcttatgaat ccatacact cttttccaca ttatctataa gttcctcttt caaaccatt 60
 ccgaaacaca ccagtatagt agacgaggag ttttatttga gcaactgcgt cttttacaag 120
 ctcccgttgc ttctgtgtt agaaaactaa cttaagagt attcaccaaa aaaaaaata 180
 aacaaacttt aagagtcata agttgcttat ggaaatatac atcctgtttt agccgttaaa 240
 aatacatgct tcgacggatc aataaattta atggtaaagc accacaattc tgtgattgat 300
 gtgggtctttt ttacgcagca agaataaaaag taccttaacc attaaatcag att 353

<210> 31130
 <211> 399

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31130

 agctnctant ttttngtttc anaacttngg nnnntctgca agnnggaaat tctgcagaaa 60
 acaaaatggt ggatcaagtg gnctcagaat aattaagaaa ggggggttga attaattatt 120
 aatgtgtcct tactaattaa aaatttaacc ttcttaatgt tactagattc aattangctt 180
 ttactactaa gttaagaaag taaagaacag aaataaaaaac ttaacccaaa gtaaaagcga 240
 taattaaaag tacatagcag aaattaaaga gtgtanggaa gaagaagaca aacacaagaa 300
 ttatactggt tcggccacaa accgtgccta catccaatcc ncaagcaacc tgctgttctt 360
 gagaattctt ttaaccttgt anaatccttt acaagccaa 399

<210> 31131
 <211> 442
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31131

 tgacaaacaa agctaatcga agcaagcaag aaataaaatc aattattggc acaaaatcaa 60
 ttgtgcaaaa ggcatttgaa atggtaacat cttttaatat tttatattca ttntcttata 120
 agttatataa taataactca tttttttatt ntgttatttt ttatatgata tatgaaagtt 180
 tggtgaaatt tatataaaga catcatacat tagattatat tatttaattt gtcttttact 240
 atatttaatt ntaatagaaa gaagattcaa aatctggtga atggccagat gctatggaaa 300
 gttggaaggt cacgcacatg agatctaata gaacttggtg cattccaaaa ggagaagaaa 360
 tcatggtaaa gaaaaaattc atttaaagtc ttgatgtcaa tattagaatt aaagacatta 420
 atggtattat gatatcatat ct 442

<210> 31132
 <211> 438
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31132

agcttgagca	tantaagtta	ctcccgcat	ntatctctag	catgcattgt	atgttggctc	60
cgtcctttgt	cacgggaagc	cggaaggctc	atatcacctt	cttaattgta	cacatggggc	120
actgcgcccc	caaatgcgca	agtaagaaga	gataattttc	cgagctctcg	tgtccgtaaa	180
atgcattcat	atcatgcac	gcataagcat	ctcttcataa	catcataatg	gacatatcct	240
gcatttgctc	gttatcatat	tccggcctca	cattttgcat	gagtcattgg	atcatcatgc	300
atatgcgttc	aacaaacttt	ntgatctgca	aaattgcata	ccatttggtt	tcatgtttgc	360
tcatccttgc	gttntcctct	acaaaacana	nacaaaaaag	ggggaagcgt	gaaacttcac	420
actacattct	tagttcca					438

```
<210>      31133
<211>      384
<212>      DNA
<213>      Glycine max

<223>      unsure at all n locations
<400>      31133
```

<210> 31135
 <211> 262
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31135

ggtacctgga gatatgtcgc gggggtcaag agaccttggg gacgtcaggt ggggtgctat 60
 tgcccaaac caagcttgac caatcccaac ccaaccggg catagtcagt caatgagaac 120
 ctatgatgta cctaaacagg cgagctcctg gcagtcaact gataaaagga acaaagaacc 180
 acanagcagg agacttgtgt ggtggctggc cagctgtgaa ctatgattga tatatgggat 240
 atgggctctg gtaatcgatt ac 262

<210> 31136
 <211> 248
 <212> DNA
 <213> Glycine max

<400> 31136

taccagctg gccttgaatc agatatccgt gcctatcgca aaggtttgtg ggttgtgctc 60
 ctttggtgac caccatacag acctttgccc ttccatgcag caacctggag caattgagca 120
 gcctgaaact tatgctgcaa atatttacia taaacctcct caacctcatc agcaaatca 180
 accatagcag aacaattatg acctcttcag caacagatat aacctggat ggaggaatca 240
 ccctaacc 248

<210> 31137
 <211> 328
 <212> DNA
 <213> Glycine max

<400> 31137

tgccaccag ctcgcccagg cgagcaaggt tgcttctcc agaagcaaca gccttctgga 60
 ggaatcttct ggagggccca agtgggctg gttgctatct gcaccttat ttttactaaa 120
 tacacccct gccttttttt tgggtgattct ttttcgtaa agttacggaa acttatgaat 180
 ttcgtaacga tacttgtttt ctttccgtaa tgttacggaa ccttgcgat tacataatca 240

tccctttttt gacttacgga atgttacgga acctactaa ttgtgcaacg atgcttcctt 300
 ttgatttccg gtgtgtcacg gaacctta 328

<210> 31138
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31138

agctttacaa ttttgtttta aatccaagcc cataaataat ataaaatcta gataagataa 60
 gataagatct agatgaaata atatctagat gagatcaaat ctagataaga taagataaga 120
 taagatctag atgaaataat atctagatga gatcaaatct aaataatata tagatgagat 180
 aaaatctaga taagataaga tctgatagaa taaaattgtc tgctcttttc aagtccaagc 240
 ccaattccgg attcaagccc aattgcttat aattctcctg aaattaaatc anaaacacaa 300
 aattagtcca gtaggtccaa ttgataaaac tgcataattan attgacaatt aagcctaata 360
 agtaattaaa atgatgacaa aaaggggttaa gaaatatgag aaaatgatga cacatcanat 420
 cccctcacac tta 433

<210> 31139
 <211> 294
 <212> DNA
 <213> Glycine max

<400> 31139

tatagcttac tgattatcca caaaaagctt cactgcatca ctatccttga ttcttaattc 60
 ttgtaataat gtgtccaacg agacagctag gcaagcactc attgtagctg gaacataactt 120
 agcttcacat gttgataaag ccactatgga ttgctttctta gaactccatg atattggtgt 180
 tgcaccatac atgaatatgt aacctataga actctttctg tcattctctgt ctctctccca 240
 atccgcatca atatatccca ctaattcctc tgagctgatg ctgtctatat ttgg 294

<210> 31140
 <211> 448
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 31140

gtttaaattg cgtttaatct tanagtttga cgtgctgac agtttttttt ttttccggag 60

ttaagaatac ggggtgtcttc caaatacaca tttttaaaaa gaatatttta ttaaatacac 120

taacgaaaat tatattttta caaattataa tttgaatata ttttaaaacc attcaaagcg 180

ctcgacagta attgctcgag aaccttcacg ggccatcaag aaacctcatt gggaaatcga 240

tgttaccctt gatagcaaga aagtgaagaa actaatttct tttagaattt tttatttatt 300

gaaaaccata caaaccacat taccgccttc ctttctaagt agcatacgtg aagcaccgtg 360

tgccacataa ctctggngtc tctactcact ctcttggtgcc tttgagttaa atcaattctc 420

acttttagtt cctttcnca ataattat 448

<210> 31141

<211> 421

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31141

tatataatga tgattatagg tggttgtttg atgccagata catctggtgc aagagttcat 60

tttatgtatc ttcttttatt atcaaattta actgaggcaa gtcattatag ttaggggtgta 120

gcagtattag catccctttt ttgagctcta gatcgggcta taaagccaga ccaaacagaa 180

atcgggtggat gtttgttgtt gctacagtca tgagcgtggg accgaattga atgtattacc 240

caaagatag atcacctatc catggaagaa gcacaagaag gactcanatt tcctcttgca 300

cgaaggtggt ctggtccaag gaccggacca aatattccca ccaattcagt gagattgata 360

cgcacatata ttgataaact acatattaat gagggtcatt nnttaatttc atgaataact 420

a 421

<210> 31142

<211> 273

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31142

attttctcaa ggaaacctgg attacctttt ctttctcggg ctcttcagca ttgctcaaag 60

aaccttctga tactttcccc ttttgtgatt gtaacataac catcttctac aattaacctg 120
 caatgtacac actgttgacc cttcaacgaa tgagctntaa cagaaagttc agtacagcgt 180
 ccatctaate tccaagctcg gagggtaaca aaacatgcac tcanaccacc aagatccang 240
 ccactggctc gaacacatnc attcaatcca aca 273

<210> 31143
 <211> 330
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31143

ttagctttga ncttatcagn caagtgtatg gaccatgtcg tagccaaagt gctcatcggg 60
 aatgggtcca agttaaactg gatgcctaag agcactttgg agaaattacc attcaatgct 120
 tcccacttaa agccgagttc aatgggtggtt cgtgcctttg acggcaccgg ccgagaggta 180
 ggggagagat cgatctccca gtacagatag gccctcacac ctgtcaagtc accttccana 240
 taatggatat taaccccccc ctacagctgt ctgttggggc gcccggtggat ccactcagtg 300
 ggagttgttc cctcaacact ccaccaaaaag 330

<210> 31144
 <211> 375
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31144

tgccaagaca atgcacacgt tctctntttt catgtccttt gacttttgaa tatatatcat 60
 ttgtctaatt agtagaatct tgggtgctnt gtaaattttg cgaactctct gtttcaacca 120
 ttttcttttt tagttcatcc tacataaata catcttataa attattatca tacatcaata 180
 tcctgaatac ttcaatatca ctaaacaaaa ctcatctcca tattagttac tccccctcac 240
 ccataaacct tctattagag aattgagcac aacaaagaaa aagtattgaa ataaaaatta 300
 caattcttac aattacaata gcagcctttt cagtaacaat gctttcatct tttcgagttc 360
 gagtgtcaat ataaa 375

<210> 31145

<211> 461
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31145

caagcttgat cttttattct atatctgaca gccaatgggt gaggcccgtc caggtagtcc 60
 cgaagaanac cggcctcatc atgataaaaa atgagaagga ggagctgatt cctattcggg 120
 tgcagaacag gtagagagtc tgcattgact ataggagggt gaaccagggt accaaaaagg 180
 accattttcc cctgccattc attgaccaga tgcttgaatg cctggcagggt aaatctcact 240
 actgtttcct tgatgggtnt tctggctata tgcaaatac tattactcct gaggatcacg 300
 acaacaccac attcaccagc cccttcggaa ctttggccta tagaaggatg cctttcggcc 360
 tgtgcaatgc ccctgggtacc ttcaagcgga gcatgattag tattttcagt gattttgtag 420
 acnattcata gaggtgttat ggatgatntc actgatatgt g 461

<210> 31146
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31146

tgctgacctt gctctagcct ctggagaaat ttttgaagga gaatgtttca agtacataaa 60
 ggaatgttnt gaaaaatggca cattgcatct cattgggcta ctgagtgatg gtggagttca 120
 ctccagactt gatcagttgc aggtgattat ttgggggttg agctgttttt ctttcattgt 180
 tattcagttt attctttcta actaactact tttgtacagt tgttgcttaa aggagttagt 240
 gagcgaggtg ttaaaagagt ccgtgtccat attcttacag atggccgtga tgttctggat 300
 ggctcaagtg tgggggtttgt ggaaaccctt tgaaaatgat cttgcaaact cgcgcgcana 360
 aggtgtcgat gctaggatag catcacgtgg aggtcgtatg aatgtcacia tggatcg 417

<210> 31147
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31147

agcttgacat ttatctggtt cgcattngnngn agcagattga tagcgtgatc tgtggatcga 60
 gatggcggaag ggtttgtaag tgggtgaaac aggcattgagt atttcgtaag taactcagaa 120
 atctcgggta agatgggtgg tgttgtagct gattgtgtgt ttgtttccac tctaataatgg 180
 aagaagggtgc tagaggggct cctatgtaga agacgacgca gttgcgaggg agacacgggt 240
 tcacctatct gctcacgttc cccctgtaac tccacaagct taccctcagt gatgaatttc 300
 atggatggag acgtgtagtc tgtcagaacc ggtcctaatag ttntgagcca ttcgactccc 360
 agaactacat ctgtgccaca taagggtagg atgtgaaagt ccaccatgaa cgtatgctcc 420
 tgcacctgt 429

<210> 31148
 <211> 243
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31148

aaacccgaaa agacctgatc tacgaattct ttgtgaaagg ttcgaagagt gtatttacgc 60
 acaggggaaag tattagcacc ccacgcgtcc gtcacaagag acgacaacct ttaatcaaat 120
 gtgcaaatat gacatcnaat tatattcntt tcccttttta cgggtcttaat gtctttttat 180
 gccttttnnta tgttttatct ttttgtggtc gacaagggtg tttccctttg cttctacgta 240
 ttc 243

<210> 31149
 <211> 464
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31149

agcttggaga ttatgcttct atgtatgana agaaagaggg agagaaagag agaggnggga 60
 gcacgacatc gaaggaagaa naaggagag aagntgaact ttgagttgtg tctcacaaga 120
 ctctcattca tcaaagttac acatgcttct atttatagac taggtagctt cttttagaag 180
 ctgtcttgag aaaacttct tgaagaagctt ctttgagaaa acttccttga gaagctagag 240
 cttagctaca cataccctc tcataactaa gctcacgtac ttgagaagct tccttaagaa 300

gattcctaaa gaagctaaag cttagctaca cacacctctc taatagctaa gttcacctcc 360
 ttgagatgag aagctagagc ttagctacac acccnctata atagctaagc tcaccncat 420
 gacannaaaa catgaanata caaaaanaaaa aagtccttac taca 464

<210> 31150
 <211> 398
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31150

tgagactnta agcagtgccct gtgattagaa tttcattnta atactttgga gtacgaaaat 60
 tcaaaaactt tataaaaaaa aaaaaaaaaa aaaaagtagc atgcagtgct acaacaaaag 120
 cacaggcaca tatgggaaaa taaatgaagt gacgtacaat taagtccttg aaagaaagaa 180
 agaaagaaaa aaaaactagt ggaagctcaa taatggagga agagaaagtg tggagcagag 240
 aaagaaacag aggtgtgtgt ggcttcttgt ggagaaggaa gaagaggaag gaggagcagg 300
 tcaatgtcaa tgtgattnta taaagctaga aaatgaatat aatacaataa ttccttacgt 360
 aagagttttt aaatgtatat tggcattaat atacttga 398

<210> 31151
 <211> 453
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31151

agcttgagga ttatggtgta cctatcacat gtggtactag gtggcggtcg ggcgatggtg 60
 cacaacaagt tttccacatc cacaatgcgc gcataaaccc accatcccct gttgcccacc 120
 tccaactgag ctacagtact cccacgtagc ccatatcctc gtttctctca acaccgggtc 180
 cccatcaatc ctcccaagct tccccaacat caaagtaata caacattcaa acagcacana 240
 ctatcacagc caagaaaaca gagcanaggc agannactct gccaaaacac caaccanaat 300
 cacagctttt ctacttaaa gaccccagta acaattcctt cgttccaatt cgttaaccgt 360
 tggatcgaac tccaaatttt actggaagtc tctagtacat aagcctacat tntgaaccgt 420
 gggatctact agcanacatc cagaactcat tct 453

<210> 31152
 <211> 186
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31152

 gcattctgtgc ggtatttcac accgcatatg gtgcactctc agtacaatct gctctgatgc 60
 cgcatagtta agccagcccc gacacccgcc aacacccgct gacgcgaacc ccttgcggnn 120
 cgatngaata taacttcnnn atatgcatgc tatacgaacg cattaccgat gagccctgac 180
 ttccccg 186

<210> 31153
 <211> 443
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31153

 agctaggctt ttgctattgc tgaagagacc agnattcaga ctttgctttg tgaattacag 60
 gtcacacaca ctacaccgct gttttttgtg acaacatgag cacagttgcc ttagctcaca 120
 acccagttct gcattccaga accaagcaca tggacctgga cttgtctttt gtcggagaaa 180
 aagttctgga gaagagaatt caagtgggtc atgttctctac tattgaatat tgatcaatat 240
 gcagacattc actaaatctc ttaccccatc taattntact ctgttttaggg acaagctcag 300
 agtggtaaca aagattttgt caaccctca agagcttgcc aggggtatta gagtagaaga 360
 gtagaattac tcctttcttt tatttcagtc tagcatagtt agcctttata gnntaactca 420
 actagtgaca gttgtaataa cag 443

<210> 31154
 <211> 441
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31154

 tgtttacacc taaatacaat catggtattn atactataaa ttaagattat attctaacat 60

tggaaaattt ttacataatt atctaataat aattggaaaa tttttacata attatctaatt 120
 cataattgga aaaattctac ataattatct agtcataatt cattatatat agcataaatt 180
 ttttgacttt taaaataatt taaacagtta ttttaattata taattaaatg ataataataa 240
 aatatttcac attgtatcag cattaacctc ctgcttccgg cttttgtgta caacatggag 300
 tctttaattt tccatcgatt atgcggctga tactttgcca cacataaatg tataagaaat 360
 atctttcaga tgttgactag tagttgattc attattttac ggggccagat tattctgaac 420
 atccattcca ctggtgcaat g 441

<210> 31155
 <211> 276
 <212> DNA
 <213> Glycine max

<400> 31155
 acaacgagat gatgcgctcc atgagagggg ggatcaaag gagaaatagag atcataatga 60
 agaagaaagg aggagaagag ggaatgatgg tgttcctaga caaaaccgaa ttgatggtat 120
 taaactcaac attcctccat ttaaaggaaa gaatgatccg gaggcctact tggagagggga 180
 gatgaaaata gagcatgttt tctcatgcaa caactatgag gaggaccaa aggtgaagct 240
 tgccgccacg gagttttccg actatgctct tgtgtg 276

<210> 31156
 <211> 290
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31156

attgagtttt ctatgccatc ccttaatgaa atatttatga tgcggtaaa gaaatgttcg 60
 atcggcgatc tgcggatgatg cttctttttt agacctcgat cggatcatctt tcttggcgga 120
 cgtcgactgg cacttttttc aatcaatata ggtagaaaat atttttttgc cgagatgggc 180
 taattgtttc gtggtcgaat aaatggaaac atgccagttt tggccgacac aaaaacgtgg 240
 ttgggctcgc acanaaaaac ctageccgacc tacattgtac attttttatg 290

<210> 31157
 <211> 536

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31157

nggcaccgtg gatgatgcan ttgcaatac nggcgaangc agctcggacc cgggatactc 60
tacagtcgac tgcaggctgc aggttgatgg tantggtgag agacaanggc atggacatgg 120
cgaaactaag tgagctccgc caattgcaca ctactgcag acttcacgaa cctanagtgc 180
cactccagaa caagactcac gtatacttgt ggtgcttacc tatctaccct agtgcacagn 240
caaccacatt gtggatcctt tgcaacggta tcacttaaac aacattggaa tgggtgatga 300
agacacttga tgataatcaa ctgatttaac tggaacctag tgtaaaacta tgcacaccat 360
taactaatat aagttatcta tgcgatggct gataagataa tcagctataa cggctcaa at 420
ctatatactg tatatatata tactatatac atatgaccgg ctaatnttgg gtgataatgt 480
gtangacaac atggacatgg taattcacgt gcggatctct cacactcagt tatacn 536

<210> 31158
<211> 529
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31158

cacacacacc actttntaa attatctgtg actaatatct acataatcct acaantaann 60
nnaaaagccg atgagcatgt gaaccatgga tnacacaact aagataccta acccgacaat 120
aagagacagt caagacgtgt atttttcacc aactatatgc acacgcatag gattacaaga 180
gcacaacatt aaaaactaca tcatgggaca caaaaatgcy acatcacaca ctgaagtacc 240
tttacattcc agccaaaact aactacctgc atgaaaagaa gagtacgaca cgcacaagga 300
gcacccccat cattgcagga aaacgacggc gaaacacaca ctgagctatg atacacctgg 360
ggagatgaga gcacgagaga aacaacagat aactcactaa gttatgatgt gagggaacaa 420
tactcagcat attacaagat gtcaaaacag agagagcaag gatatgactc cctctaccgg 480
aaaatatggc gggcaaaagg ccataggcct actcacaacc gacaccccc 529

<210> 31159
<211> 439

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31159

 agcttgngtg tcntgtttat aacttcacaa gagatatttg aaaaaggat tacaaccatt 60
 tctctatctc tttctcttat ttntatatcc ttatgttttc atatgataag aagaanaata 120
 aattgaatta agaaaaaaaa ttgatcataa tgatttatcc ctttaggtat aatacaatac 180
 caaaattgaa atatttatag catattaggt ctacaatttt tattcttaca atcttaaaaa 240
 ataatcattc tcattatgta tgtcttccta gatccaatat gcaataaata tatcaatttt 300
 agccttccac catatctaaa ggaataaacc atntaataat aatgactatg caattatatg 360
 tgaatgataa aattagtttg tgggtacata atcacttata naaatcatac anttttatat 420
 tttaatttaa cctataaat 439

<210> 31160
 <211> 376
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31160

 tccattntcc tttctacttt gaatgagtaa ttcttaagtt tcatacatca tcagggtccat 60
 ccttatattt ttttttcctt ctaattatta aataaaaaata ttgttttctt gaagtctctt 120
 ccttccatgt aataatgaaa ccgtaagatg accaaattaa ttntacttgc tagttatttg 180
 aagaaccaac gtcgcacgcc aaaagtcaaa acctacaacc catctgtcat cccgattcat 240
 tactctcggg attacttcag ggtcatttca ttgttatctc ttcttccttt tcacaccctc 300
 atttaagatt taaaacacct aanataccct tctttctctt taagaattaa aatcttctct 360
 cctactcatg tccatg 376

<210> 31161
 <211> 435
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31161

gctttcatct agtgtttatc agagcacaag agcttcaagt aggtgctcct tanacctcca 60
 ttaattnttt tctttacctt ctcttcatt gttgtttctt catttttctc catgtatctc 120
 ctcacatgtc ttgtgctaaa ttttgtaac atgattcttt agagtttcca ccgattaaac 180
 ttgctataga agctagattt gattntctat tgttcaaatt tcttgttctt gttcttgaac 240
 catgaattgt gttgactnta ngttcctttg agttttgtct tgttattttt tgtggctgat 300
 acctaaacca tanaattctt acaaaaaat taaattagaa gaaaacctan aaaatctaga 360
 gtgacttggt cacctattgt agtntgtca tagaagtcac gtctagtcac gaaacttgtc 420
 acataagatt tctta 435

<210> 31162
 <211> 314
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31162

tggtgccatt gaactcatgt gctacaacca agccatgttt gggtttgcac cctatgggtcc 60
 ttattggcgc caactaaaaa agattgtaaa cttagaaatc ctctccaatc gccgagtaga 120
 gcaactacag cacgttcatt tctcagaagt tcaaagttca atcaaagagc tcttcaatgt 180
 ttgggtcaagc aaaaagaatg agtctggcta tgcgttggtg gagttgaatc aatgggtntc 240
 tcatttgaca ttcaacacgg ttcttcgagt ggtcgttgga aagcgacttt tcngtgctac 300
 aactatgaat gatg 314

<210> 31163
 <211> 443
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31163

agcttgagct aaatttgact accatnacct tgaccagggtg agaatgccaa tcttcctcgt 60
 gaagcaaaaa aaaaaaaga agagaggaaa atttccaatc aaaggaaaaa ggagaaagaa 120
 aatttccaat caaagaggaa gcaaaaaaag gagagaagga nnaatttcaa tcaaaggaaa 180
 aaagagagga aaggaaattc ccaatcaaag agtgggagaa agcaaaaaga aaagaaagaa 240

aaattcccaa tcaaagaatg ggagaaagaa aaaaagaaga aagctcctgg tcaaagaaac 300
 cagaagatat gtgccgagag gtccttggac cagacgatat ccgaacaata cagaattgtc 360
 accaaatgaa caaaagaaag aaagggaaac catgacctan aagtgggtctt ctccctttat 420
 taccaaccaa aatcctgtgt gct 443

<210> 31164
 <211> 300
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31164

gtgaagtgaa gaatatttat tactgagaaa cttaagtgtt cattatagtg gaatcttctt 60
 tcctaataat ttttttttct ataacgctgt atcttagtta aatgatcaaa gtttaattct 120
 acttcaacta atctaaatgt ggtaataacg tccttttggc tgcattgcctt gttatatagg 180
 gggaaactat ctaanatgaa acttaatctt attanggagg tatttttcagc anaatccaaa 240
 ctcatctcta ctttatctta ttgcttatgt tncagggcac accactggat tctatactca 300

<210> 31165
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31165

agcttnccct tatgaatctc aagtgcnctg gcacgcctat tattctcctc ttaaacacgt 60
 gtacgttgag catttctcca ttgtccaagc atatactata tgaactatga aatcagttct 120
 atatgtggag ataataaaat gacatgactt ttaatgtatt ttacagtga tgaataaatc 180
 attcatatta attatgtaga agctagaaaag gataaaatga tacactntct tcttcttctt 240
 ttctctttta atttaaacaa ctaanagaat tatcattntt tttattttca ttntcttnt 300
 tatccaaaca tgacatagaa tggtttagtt agaanaatat tagcaaaaca canacagcgg 360
 ngctgaagtg aattagttaa gggctctttn tagtccaagg accggcgctg atgcatggaa 420
 tatgaaaata tatatataaa acgaattatg ataaaac 457

<210> 31166

<211> 392
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31166

agcttgtcca ttaaaatatg tntttgaagt tagtcatttc aatttctgac taagtaaaat 60
 ggatcatttt taagggtccaa cgccttgaaa tgatcacctc ttaagtaaaa aaaaaaatca 120
 cttgataagc tagaactacg taggtctgat ttcttcatcg caattgagga tacgtaggag 180
 caaaagcccc gcttttgtcg accaccccgga gagatcgta atgggtccaac gccttaacgt 240
 ttctctcctt tcaaaatcaa aagatcattt aatgggtccaa caccttanat gacctttntg 300
 ttcaatcaaa atatatcttg caaaaagata aaaaacaact taaccaaaaca ctntgttccg 360
 aaagaactac gtangtcttg attcctcatc gc 392

<210> 31167
 <211> 385
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31167

ttctaaatga taggctcaga atgcagaaga agtagcaatc aatttaataa tgttctttat 60
 acatgcaaga caaaattgat tgcaataata aatgagataa gggaagagag aaatataaac 120
 tcgatttata ctgggttcggc cactccccgt gcctacgtct agttctcaag caaccactt 180
 gagattntcc tttctctttg taaaaccctt ttacaaagtt tgaaccacac agggaacaacc 240
 catcccttgt gttcagaaat tcttacaact taagagacce tcagtctctt aatcaatctc 300
 tttgattaag aagaagaaga agaagaattc tctcttttaa gagaaagata atacaatgaa 360
 gttccataaa ctcttaatatg atttg 385

<210> 31168
 <211> 338
 <212> DNA
 <213> Glycine max

<400> 31168

tctgttctca attacgaacg tctcgatata ttacgggaca caatcggaca cccgagttaa 60

aagttattgt cgtttgaatt tgctcagagc ttctattgtc aattacgagc gtctcgatat 120
 attacgggac tcaatcggac atccgagtaa aaagttattg tcgtctgaat ttgctcaaag 180
 cttctgtttt caattacgag cgtcctgata tattacgtga ctcaatcgga catccgagtc 240
 aaaagttatt gtcggttgaa ttgctcaga gcttctgttt tcaattacga gcgtctccat 300
 gtattacgag actcaatcgg acatccgagt aaaaatta 338

<210> 31169
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31169

aaatgtttca aacttttatg gtcagtgtaa accacaaagg gcctgcccac cagatagtgc 60
 ctccagcgtg gaatagctaa cactaaagcc atgagctcct ttccatacgc agattttgat 120
 aaatcgccct ctgatcaagc tatgctaact aaagctatag gctgcctctg ctgcatcaaa 180
 acagcaccaa ttctctccc cgccgcatca cattccactt caaacagaat agagaaatca 240
 ggtaacacta gtactggagc tatagtcatg atctgcttca gatgattgan agcctccaga 300
 gcatcttttc cccaaataaa gttattcttc ttagtcattt cagtcaacgg gttagcaatt 360
 gtaccataat cctttgataa tttctgtata accc 394

<210> 31170
 <211> 420
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31170

tccacttgta tactgcaacc ataataaaag ctntctattn ttccctcccg tggatgtagc 60
 ctttctcaag gtgaaccacg taatcctgtg tgtgttcttt ctcatctctt tctctttcaa 120
 ttctgatgca atttgtgtgt atgataattt tggtatgctg catctattgc tgctgttctt 180
 gcttggttctt catcacttcc ataatagcct tggctatcga tgtaggcacc ttgggcaccg 240
 gtgagggccg tatgaactgc taacgttccc tttntactca ttatttattt cttntattg 300
 gtaatttatt caaatgttct catcgtcacc tttctccttt cgctatgtn ttttctttt 360

ggccaactat ggcgaaactg catcgtgtcg cegtgttgc gccaccatct tcgcgatcgt 420

<210> 31171
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31171

gcttangtgt tttatgacca atctaaattc tattaaatac atatattcgt agcatgtgtt 60
ttcaaggaag tctacgttac attaactttg attatatact aacttgattg gttatgtcat 120
gtttgtttta taagatttga tgcccaagtt ttctgaagtt gttgaaaatg attcacaatg 180
tggtgaaaaa agaagcaact aatttatcaa aggattntga agagttaacc cctactaaac 240
acaatttcag ttgctattca gctatgggag aattgactcc taatgaaagt tagtctgcaa 300
cccanaataa ggataaatga tcttttgctc aataattctg gagatttgat tcccgttgga 360
actgtatttg gaactcanat acattcacag cccgaanaca ttgaaggtgc atataagcaa 420
ctgggtggat gtttgataat gtcanaacta at 452

<210> 31172
<211> 323
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31172

ccaaacgcac ctttcttcaa acattcggtc gtaggacttg ctataacgct aacattctgg 60
ataaagcggc gattacatga tgcaagacca aggaaagatc tcacgctacg aactggtaga 120
aggctcggcc aagtcttgat agcatgcact tttgtttgat caacggatac tccatcttta 180
gacaccacat atccaagnac accacacttt caaccaagaa agcacactnt tccctcttgc 240
catagagtnt tcgggctctt aggggtctcaa atatttggtt canatgagtg aaatgcccct 300
ctatagatnt gctatacacc aat 323

<210> 31173
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31173

agctttacat tactcaccca cctatggnta gcaaagtgtt ttgtgtggga atttagatga 60
tcattttgtg ttttccttaa gttggacaat gaggcgagtc gaggtggctt tggcccccttg 120
gccccgtaaa aggttgagtg agtttaggtt atatttatcc ttgccctgct aaggtgcttc 180
attgtcatgc tggggtagtt nttcatctta ctcaagtact tcttccttat gctaaggtag 240
cttttcctct cagtgttgag ctacctcctt gtctatgtca gttccctcat cctcaaactc 300
aacataacct tattgagttg atttcattnt caccctaana aagttgactt ggattgngca 360
tcatttaatg ctattgtan gtggctctgc ttgangtcgt ggagatggta gtgtanaatc 420
tctatgatga ca 432

<210> 31174
<211> 343
<212> DNA
<213> Glycine max

<400> 31174
tgtcctcggg gacgaagaca attgaataag ccccttcgag cttttcacag gcgtcaacga 60
ttctcaaaat gaagggctctg tgtttggaag tggcgatgag gtggagaaca acctcggtgt 120
cggaggtagt gttgaagatg gaccgcgtgt cctcgagggtt ggttcggagg gtgcggtagt 180
tgacgaggtt gccgttggtg gccacgccga cggagccgaa gcggtagccg gcaacgaagg 240
gttgcacgtt tttgagcatg gattggccgg cggaggagta gcggacgtgg ccgatggcga 300
ggctgccggg gagctgggtc agcttcgact ggttgaacac gtc 343

<210> 31175
<211> 443
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31175

agtttttaaat tttatatnta aatntctaaa agctgataca aatagtntaa acttctgcta 60
atcgattaca taccttgtgt aatcgattac aggcttttaa attcaaattc aaaattttca 120
aattttttca gaaatcaact tagccactgg taatcgatta catcatctgc taatcaatta 180

ccagagagga aatatcatat ttttgaaaag ataattgttc tttaaaaaac ttttgtaaaa 240
tatttccttt agccaaacct gtgcaacatc aattaaggaa ttctttctaa gattctaact 300
atgtatatcg ttcttcttgc atttctgaat tcttgactta aatcgcgctt atctttggca 360
tcatcaaaac ttcatatcat atatgcttct acatcctana gtaatacttt gaaagacaga 420
gaagacatca naatgatttt tca 443

<210> 31176
<211> 245
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31176

tcattaagaa gcttactcca gaagcttcct cgtggcttct ttgagaagct ttctcaagag 60
acttctttga gaagctagat ccttatctat ccacaccct ctattaacta aattaacctc 120
cttaaaaata attacggata aaaataacac aacaaataat tcaacatcaa acataattac 180
taataattta tatatatata tatatatata tatatatata tatatatcan ggtgttacac 240
ctact 245

<210> 31177
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31177

tctgctntgg attatgatnn ctatacaaaa gttagtgtga taaagtgact aacggngacc 60
cactctgcat gtctgtcaat tgaccatgcy atatccacgt cgtgtaattc ttcttaatcc 120
catcacacaa tagatggtcc catatgtcgt ccagtatttg tgccttccg ttcaaacaat 180
tgatacgagg aactaatat ttccatctt catccagtcg acttctttnt gaagtaaatt 240
gcaagaactg ctcgacgcct tccatcatatt ctgggttgat gtgactntca ttcatcaaac 300
ttcgatecat ctcagtaata actctgtgat actcanagtt attcgatgct tganaatctc 360
actnntttat tatagggtgtg gccctatccc attcangaag accgtctntt atggtagctt 420
catacgtca 429

<210> 31178
 <211> 370
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31178

 taagtctaga ttaatttaat tgtcactatc catggcaacc tctttgataa tttttttttt 60
 cttttttaag aggaaacaga tcgagacact gacgaataca aatcacaaaa ccgtgaagaa 120
 aattcacaaa aacgctaaaa tttcataagt tctcaaccca cattccccc aaacccacaag 180
 ttttcttcatt tttctcagca aacaagcagg aaaaaaaaaa ggcaaatcag gaggattgca 240
 cattatgcac aaagtttagat ctgagaaaaa aaaaaaccca aatgcatgca aaaaagaata 300
 aagaaataaa caagttgaat caacaatgat gaaatntgaa aataaaaacta aaaaaaaaaa 360
 agtagaaaga 370

<210> 31179
 <211> 450
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 31179

 tatgctttta tttatatgga cttaccttga attaatcctt ttgatagccc ttttgagcct 60
 tgtttccctt tccttggttt gaagctcact acaagcctta agtgaaaaac catgatatta 120
 ccatatcctt aaggaatttt ggagcttttg aattgttttg ggaataagtg tgggggggttt 180
 ttgtttcatt ggacaacttg ttttggtgac tatgcttcat gatgtatttt gggccatact 240
 tgatgtacat tgtatattgg ttaaagtgtg gacatgctga atgaaatgtt gtttctcaaa 300
 ggcaaaaaaa aaaaaaaaaa aaaagcaata aagttgagtg aataagatct ttaatggcac 360
 aagaatgatg aaactcttga gtctactctt catgggtaat tnttatcttt acttcttttt 420
 tntttttctt aatatgcact tattccccctt 450

<210> 31180
 <211> 334
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31180

ttagctttat ctttataatc ttgcatgat gatgatctgg cctctgtgca ngctctgtaa 60
 catagcaaat aaagtctcaa agctgaatga ggaaggtcct gtagaaacta gagtaggaac 120
 tggcaaagga gtgaatgctg aagctagaac tggtaaagat gaagctggaa tagatggagc 180
 tgcagtagaa ggagctactg atggnggaat ctcaaaagat gggagagtct ttgacctttt 240
 ccccttggcc ttgcgaggcc ctctaaaagt gattgagggg tcatcgacgt tccaccaat 300
 tttcttcatg taagccaaat taatggttgg actg 334

<210> 31181
 <211> 335
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31181

tgagttnttt taaaaatcat atttagctaa ataggagaga ccatgaactt taaaaaaac 60
 ctattaagtc tgatgaaccg acctgttttag caataatatt ntatattaaa gatattatta 120
 ttaatatgat atatagtata attattatat ttaaattata aaattatttt aagagtttga 180
 caattataat tagtgcttga aatatcttaa ttctgtataaa tataaatgtg tacaaaaata 240
 tacattcttt tctttgggtg agacttaaaa gacttttagca atatgtcacc cacaatggtt 300
 ttccatattt tattgttaaa attgtcttat tattt 335

<210> 31182
 <211> 452
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31182

ggtgttgaac ccatcattag tgaccttata atcccagcct gaggtaactt atgcttanta 60
 acggtaccca actgcgttgt atagaaatct atacctggcg caaaaggcta tgggttatgc 120
 ttctctggcc aacaccacac aaaacttttt ctttcatgc cgcaacctgg agccattgag 180
 caacttggaa cttatgctgc aaacatttac aacaaacctt ctcaacctta gcaggcaaat 240
 caaccaccgc agaacaatta tgacctcttc aagcacaaaa tccattcccg atggaggaat 300

aacctaattct tagaggtcta gccctaacaa caacacagca gcctgctctt tctttcaaat 360
 gatgctgcta aacaagcatt cattcttcac aatcaacaca gcacagccca gaacacaaca 420
 gttgagctct cgaaccttct cgagactgta gg 452

<210> 31183
 <211> 236
 <212> DNA
 <213> Glycine max

<400> 31183

tctatagaag gttcgttcct aatttctcta caattgcac acctctcaat gagctggtga 60
 agaagaatgt ggcatttacc tggggtgaaa aacaagagca agtctttgct ttgctcaaag 120
 aaaagcttac taaggcacct gttctagctc ttctgacta ttctaagact tttgagctag 180
 aatgtgatgc ctctggagtg ggagttggaa ctgtattgtt acaaggtggg caccct 236

<210> 31184
 <211> 512
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 31184

ctgactatga ntcnattgca naccgggaac cgagcacgac caggcacctg taaccgccgc 60
 cggcggttta ctttatcttt atcannccg ccagacgaag accagggctc tgagcaagct 120
 ctgcaacaca ccagananag ncgcaaagca gaaggaggaa cgggctgcag aaactaaacg 180
 acgaactggc aaaggagtga acgctgaagc tcgaactggg caagatcaag ctggaataga 240
 tggagctgca atagaacgag ctactgatgg gggaatctca taagatggga gagtctttga 300
 ccttttacc ttgggcttgc gaggccctct aaaagcgact gaagggcacc gacgggccac 360
 cacattttct tcatgaacgc caagtaatgc gtggctgaga ctcccacaag ataatgaatt 420
 cgaactggca tcctatgctc ttgctaaagg agcaatgata gcagagaaga ctatcctccg 480
 gagtattctg tgctacacac aaactaacag ag 512

<210> 31185
 <211> 411
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31185

tataagaaca aaattgccta aatcatttcc aaatatgcat gtgaattagg aagcatcaac 60
aagaattaag ccaaggctat tgtgcaagca atcaatgggg caaaaaaacac taaaagatta 120
tgatgatgga tggctcaa at tctcaciaag gtaaacttat cactttcaaa ttgagctttc 180
aaaactatca tgacatgtag aggaaaaaca aggatttcaa atcacaaaat gtcaagagac 240
ttttattttc agaacaatta cccattactt gaacatatcc tataattcan agacaaacat 300
gcaaatttaa cacaacaaaa ctaacaaaat taaactaatt taacacaact aacaaaacca 360
aaaccaaaga acacactccc ccccccata cttaaacaac acattgtcct c 411

<210> 31186

<211> 455

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31186

agctngntat ttacggatga naaganaacc gaaagtgaac gaaataaaga tgaaagccaa 60
caaaacaaga aatgaattga aagtctcgga ttcgaaaact tatcggttga agaacgaaga 120
acggtgaaga acgacaaaaa atcttcatga aattgctcac gaaaatgtct cggaagtgtt 180
acggaagcac ctgggcttgg attttcttca cgaaaacatg gtttttcacc caaacagtt 240
gaaatgcata gccaaagggg ttaggggccc tttggaacag ccccccttg cctatttata 300
agaaaaaggg gaggaggttg ccgcctagca ngcccagggtg agctgagttg cttcctcctt 360
aagtaaccaa gcttccanaa ttcgaaaaat tgaaaatggc tattngcacc cncatcttga 420
taagtcaccc ccttttcgta attacgaaaa agtat 455

<210> 31187

<211> 401

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 31187

tgtcttataa gctaaaatct acttatatac ttaactttta tagatatacct ctcatctaac 60

atgcttaatt taggacncaa agacctttca agctgagatg aactgactaa tgcctccatg 60
 tgccacagaa tccagacata aactatcatc tcttaacgcc tcactcttat tatcatcatc 120
 acaccacata ttaactttct caccgtaggt gaactctaca tagcatctct cacagttgtc 180
 catgggctat tcttgcttat aaacatctct aacatgatca cataaacctt aaacaaaatg 240
 ggatgtctac tca 253

<210> 33624
 <211> 489
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 33624

ncgccgcccg cgtgagnctt atttgatacc tttgaattcc gcgacctatg aatactcagc 60
 tcgtccctcg tcgggactat gtttccactt gctaattctca accaccgcaa tgacatatta 120
 cacatttgac gcatgataca gtgcaaaact ccttagaatg ctagtaattg aaatcatctg 180
 ggcataggat ggaattactc tattgagtgc gacatgcctg atgaccatag tgaatgtatg 240
 aatacatgca ttctgacgat gccacacaat aatctaacaa agtctgcttc tttactacct 300
 ccctaagagt ccatgatacg ctgctacat attaataccc gatgcttata aatcctcatt 360
 gatcaggatc tatcaagatc tgcatacct caatacaagg tgcataagct ctacgctctg 420
 taatcatacc atgttggcac tgatacccct gcagttttat aataaattac agtgctttac 480
 atcattacg 489

<210> 33625
 <211> 427
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 33625

ttcttggttc gctatccgca gactcaacag aagtcagttg ggaaagaaat cagactatgt 60
 gcacgaatct tatacccgat gggttattga taggaccaag agctttggcc taccctaccg 120
 cttaccocaga tacctatcgt ccaccatccc accatcatcc ttgcctatcc cttttgatac 180
 taaggaagag tttcatgaac aattaaccaa agaaaggcaa gagaaagaaa cttggaagag 240